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TITLE OF THE INVENTION (500 characters maximum)

STEM CELL POPULATIONS AND METHODS OF USE

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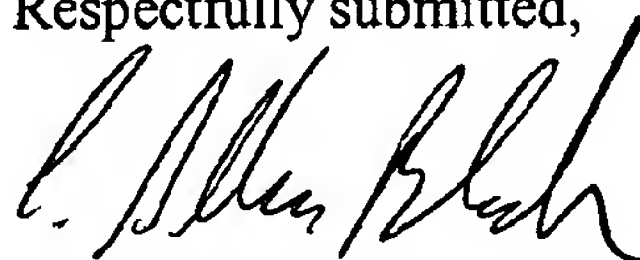
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STEM CELL POPULATIONS AND METHODS OF USE

FIELD OF THE INVENTION

The present invention relates to populations of stem cells, methods for isolating these stem cell populations, and methods of reconstituting, repairing, and regenerating tissue using the same. The invention additionally relates to methods of screening agents that promote growth, engraftment, and differentiation of stem cells.

BACKGROUND OF THE INVENTION

Stem and progenitor cells (SPC) reproduce and maintain developmental potential until specific biological signals induce the cells to differentiate into a specific cell type or tissue type. Adult stem and progenitor cells (ASPC) are small populations of SPC that remain in tissues of an organism following birth and are continuously renewed during a lifetime. *In vitro* colony assays have demonstrated that bone marrow (BM), mobilized peripheral blood (MPB), and umbilical cord blood (UCB), all contain a variety of ASPC. Bone marrow is particularly rich in multipotential ASPC.

ASPC populations that give rise to a lineage are likely to be heterogeneous. Thus, not all stem cells are CD34⁺. Similarly, while many common lymphoid progenitors are CD7⁺ and CD3⁻, some are CD34⁺ and others are CD34⁻. In part, such heterogeneity may reflect the fact that cell surface antigen expression, including CD34 expression, can depend on cell cycle or activation as well as developmental potential.

Some studies have suggested that the most primitive human hematopoietic stem cells (HSC) express the CD34 surface marker (i.e., are CD34⁺), lack obvious lineage commitment markers (designated Lin⁻), and express low to undetectable levels of other cell surface markers including CD38, CD71, CD45RA, and Thy-1. See, for example, Terstappen *et al.* (1991) *Blood* 77:1218; Landsdorp *et al.* (1993) *J. Exp. Med.* 178:787; Cicuttini *et al.* (1994) *Growth Factors* 10:127; De Bruyn *et al.* (1995) *Stem Cells* 13:281;

Di Giusto *et al.* (1994) *Blood* 84:421(1994); Hao *et al.* (1995) *Blood* 86:374; Huang *et al.* (1994) *Blood* 83:1515; Muench *et al.* (1994) *Blood* 83:3170; and Rusten *et al.* (1994) *Blood* 84:1473. Conversely, other studies using long-term murine bone marrow transplant models have indicated that CD34^{lo/-} cells contain a hematopoietic stem cell population that is capable of durably generating lymphoid and myeloid lineages following transplantation. See, for example, Osawa *et al.* (1996) *Science* 273:242; Morel *et al.* (1996) *Blood* 88:629a; and Jones *et al.* (1996) *Blood* 88:487, in which a population of small Lin⁻ CD34^{lo/-} ALDH⁺ cells capable of durably generating lymphoid and myeloid lineages following engraftment was identified.

Some evidence also suggests that ASPC that appear to be committed to a certain cell lineage, such as blood cells, may retain the ability to form additional tissues, such as muscle or nerves, under appropriate conditions. For example, CD133⁺ mesenchymal stem-like cells generally express more neuronal cell markers than CD34⁺/CD133⁻ cells (Padovan *et al.* (2003) *Cell Transplantation* 12:839). Additionally, SPC become progressively restricted in their developmental potential. Thus, human cell populations that express CD45 and CD34, but not CD38, are highly enriched for multipotential (pluripotent) hematopoietic stem and progenitor cells; whereas, cells that express CD45, CD34, and CD38 concomitantly are more restricted developmentally. Similarly, human stem cells that generate endothelial, but not hematopoietic, colonies *in vitro* generally express CD31, but not CD45 or CD34. Mesenchymal stem cells generally express CD105 and CD135. Chen *et al.* (2002) *Proc. Natl. Acad. Sci. U S A.* 99:15468; Chen *et al.* (2003) *Immunity* 19:525; Pierelli *et al.* (2001) *Leuk. Lymphoma.* 42:1195. Morita *et al.* (2003) *Eur. J. Haematol.* 71:351.

Aldehyde dehydrogenase (ALDH) is a marker that can be used to enrich APSC. See, U.S. Patent No. 6,537,807, herein incorporated by reference in its entirety. A fluorescent ALDH reaction product must be used to identify cells via flow cytometry because the marker is not expressed on the cell surface. See, for example, U.S. Patent No. 6,627,759, herein incorporated by reference in its entirety. Expression of ALDH is elevated significantly in hematopoietic, neural, and potentially other types of ASPC. Cells can be further enriched by gating on low granularity, i.e., side scatter channel (SSC^{lo}) cells. ALDH-positive cells do not co-segregate with CD34. CD34⁺ cell

populations include ALDH^{bright} (ALDH^{br}) and ALDH^{dim} cells that, respectively, express high and low levels of enzyme (Storms *et al.* (1999) *Proc. Natl. Acad. Sci. USA* 96:9118). The ALDH^{br} cells include virtually all of the stem cells, as evidenced by this cell population's ability to generate multipotential cell colonies *in vitro*, its ability to reconstitute NOD-SCID mice over a long term, and its ability to rapidly home to the bone marrow in NOD-SCID mice. Conversely, the ALDH^{dim} populations, despite being CD34⁺ cells, have very limited colony-forming ability, fail to home effectively, and only generate short-term reconstitution in NOD-SCID animals. Thus, ALDH expression can be used to distinguish and isolate functionally active from functionally inactive ASPC CD34⁺ cells. Heterogeneity in umbilical cord blood (UCB) ALDH^{br} populations with regard to CD45 and CD31 expression has also been reported (Hess *et al.* (2003) *Blood*, 102:383A).

Technologies for isolating and preparing therapeutically active ASPC from bone marrow (BM) or mobilized peripheral blood (MPB) are particularly useful because the patient only needs minimally invasive procedures for stem cell therapy. Moreover, because ASPC populations are syngeneic, the grafts will not be subject to rejection. Allogeneic ASPC populations derived from the BM, MPB, or from umbilical cord blood (UCB) of graft donors are also useful. However, to prevent graft rejection, histocompatible donors and immunosuppressive protocols that do not interfere with graft function are needed.

The therapeutic utility of ASPC is well established, and considerable evidence exists that ASPC cell populations can also generate non-hematopoietic tissues in transplant recipients. See, for example, Verfaillie (2002) *Trends in Cell Biol.* 12:502; Ferrari *et al.* (1998) *Science* 279:1528; Gussoni *et al.* (1999) *Nature* 401:390-394; Orlic *et al.* (2001) *Nature* 410:701-705; Jackson *et al.* (2001) *J. Clin. Invest.* 107:1395-1402; Grant *et al.* (2002) *Nat. Med.* 8:607-602; Mezey *et al.* (2001) *Science* 290:1779-1782; Brazelton *et al.* (2000) *Science* 290:1775-1779; Krause *et al.* (2001) *Cell* 105:369-377; Petersen *et al.* (1999) *Science* 284:1168-1170; Lagasse *et al.* (2000) *Nat. Med.* 6:1229-1234; Rehman *et al.* (2003) *Circulation* 107:1164-1169. In addition, some evidence suggests that transplanted ASPC induce host stem cells to repair tissues (Verfaillie (2002) *Trends in Cell Biol.* 12:502). However, because cell populations from stem cell sources

contain only a small percentage of ASPC, there is a need for methods of identifying the functional minority versus the non-functional majority. Once these cells are identified, therapies can be improved using custom-engineered grafts that not only contain all necessary cells for a therapeutic result, but also lack potentially dangerous, contaminating cells. Moreover, identification methods allow the useful ASPC to be concentrated, which reduces the amount of material that must be transplanted, thereby reducing tissue damage and toxicity and increasing efficacy. In addition, such selected cells can be used to generate mesenchymal cells that can be used to repair or replace tissues such as nerves, muscles, and endothelium. Stem cells can also be propagated *in vitro* and expanded into mesenchymal and hematopoietic cell lines to further increase the number of SPC or tissue cells that can be used for transplantation.

SUMMARY OF THE INVENTION

Populations of stem cells and methods for their isolation and use are provided. These stem cell populations comprise aldehyde dehydrogenase positive (ALDH^{br}) cells isolated from bone marrow, and ALDH^{br} CD105⁺ cells derived from any stem cell source. These populations can also be enriched for cells expressing such surface markers as CD34, CD38, CD41, CD45, CD105, CD117, and HLA-DR, and/or negative for such cell surface markers as CD3, CD7, CD10, CD13, CD14, CD19, CD33, CD35, CD56, CD90, CD127, CD133, CD135, CD138, and glycophorin A. The stem cell populations of the invention are isolated from a stem cell source such as bone marrow, peripheral blood, umbilical cord blood, and fetal liver.

Methods of the invention comprise isolating and purifying stem cell populations from stem cell sources, and methods of using these cells to reconstitute, repair, and regenerate tissues.

EMBODIMENTS OF THE INVENTION

What is embodied is:

1. A cell population of ALDH^{br}CD105⁺ stem cells, wherein said cells are capable of multilineage development.
2. The cell population of embodiment 1, wherein said stem cells are derived from bone marrow.
3. The cell population of embodiment 1, wherein greater than about 10% of the cells within said population also express at least one cell surface marker selected from the group consisting of CD34, CD38, CD41, CD45, CD117, HLA-DR, and combinations thereof, wherein said cells are capable of multilineage development.
4. The cell population of 1, wherein said population does not substantially express cell surface markers selected from the group consisting of CD3, CD7, CD10, CD13, CD14, CD19, CD33, CD35, CD56, CD90, CD127, CD133, CD 135, CD138, glycophorin A, and combinations thereof.
5. The cell population of embodiment 1, wherein greater than about 10% of the cells within the population are side scatter channel low (SSC^{lo}).
6. The cell population of embodiment 1, wherein greater than about 10% of the cells within the population are side scatter channel high (SSC^{hi}).
7. The cell population of embodiment 1, wherein said population is capable of engrafting a mammal.
8. The cell population of embodiment 7, wherein said population is capable of engrafting hematopoietic cells.

9. The cell population of embodiment 7, wherein said population is capable of engrafting a SCID/NOD mouse spleen with human B cell precursors.

10. The cell population of embodiment 7, wherein said population is capable of engrafting human thymus tissue transplanted into SCID/hu Thy mice with T cell precursors.

11. The cell population of embodiment 10, wherein said T cell precursors are capable of developing into T cells expressing CD4 or CD8.

12. The cell population of embodiment 7, wherein said population is capable of engrafting mesenchymal cells.

13. The cell population of embodiment 12, wherein said population is capable of engrafting tissue selected from the group consisting of bone marrow stroma, bone, cartilage, tendon, fat, smooth muscle, cardiac muscle, skeletal muscle, nerves, oligodendrocytes, fibroblasts, endothelium, and combinations thereof.

14. A composition comprising the cell population of embodiment 1 in a pharmacologically acceptable carrier.

15. A method of reconstituting blood tissue in a patient in need thereof, said method comprising introducing the cell population of embodiment 1 into said patient, wherein said cells are capable of engraftment.

16. The method of embodiment 15, wherein said patient is in need of treatment for bone marrow ablation.

17. The method of embodiment 15, wherein said patient is in need of treatment for cancer.

18. The method of embodiment 15, wherein said patient is in need of treatment for sequelae related to cancer therapy.

19. The method of embodiment 15, wherein greater than about 10% of the cells within said population express a cell surface marker selected from the group consisting of CD34, CD38, CD41, CD45, CD117, HLA-DR, and combinations thereof, wherein said cells are capable of multilineage development.

20. The method of embodiment 15, wherein said population does not substantially express cell surface markers selected from the group consisting of CD3, CD7, CD10, CD13, CD14, CD19, CD33, CD35, CD56, CD90, CD127, CD133, CD 135, CD138, glycophorin A, and combinations thereof.

21. The method of embodiment 15, wherein greater than about 10% of the cells within said population are side scatter channel low (SSC^{lo}).

22. The method of embodiment 15, wherein greater than about 10% of the cells within said population are side scatter channel high (SSC^{hi}).

23. A method of repairing or regenerating a mesenchymal tissue in a patient in need thereof, said method comprising introducing the cell population of embodiment 1 into said patient.

24. The method of embodiment 23, wherein said mesenchymal tissue is selected from the group consisting of bone, cartilage, fat, endothelium, muscle, and combinations thereof.

25. The method of embodiment 24, wherein said cell population of embodiment 1 promotes neovascularization.

26. The method of embodiment 23, wherein said population is introduced to correct a bone defect.

27. The method of embodiment 23, wherein said population is introduced to correct a cartilage defect.

28. The method of embodiment 23, wherein greater than about 10% of the cells within said population express a cell surface marker selected from the group consisting of CD34, CD38, CD41, CD45, CD117, HLA-DR, and combinations thereof, wherein said cells are capable of multilineage development.

29. The method of embodiment 23, wherein said population does not substantially express cell surface markers selected from the group consisting of CD3, CD7, CD10, CD13, CD14, CD19, CD33, CD35, CD56, CD90, CD127, CD133, CD 135, CD138, glycophorin A, and combinations thereof.

30. The method of embodiment 23, wherein greater than about 10% of the cells within said population are side scatter channel low (SSC^{lo}).

31. The method of embodiment 23, wherein greater than about 10% of the cells within said population are side scatter channel high (SSC^{hi}).

32. A method of inducing immunological tolerance in a patient in need thereof, said method comprising introducing said cell population of embodiment 1 into said patient, wherein said cells are capable of downregulating alloantigen recognition and response.

33. The method of embodiment 32, wherein said population is introduced to prevent graft versus host disease.

34. The method of embodiment 32, wherein said population is introduced to ameliorate graft versus host disease.

35. The method of embodiment 32, wherein greater than about 10% of the cells within said population express a cell surface marker selected from the group consisting of CD34, CD38, CD41, CD45, CD117, HLA-DR, and combinations thereof, wherein said cells are capable of multilineage development.

36. The method of embodiment 32, wherein said population does not substantially express cell surface markers selected from the group consisting of CD3, CD7, CD10, CD13, CD14, CD19, CD33, CD35, CD56, CD90, CD127, CD133, CD 135, CD138, glycophorin A, and combinations thereof.

37. The method of embodiment 32, wherein greater than about 10% of the cells within said population are side scatter channel low (SSC^{lo}).

38. The method of embodiment 32, wherein greater than about 10% of the cells within said population are side scatter high (SSC^{hi}).

39. A method of producing neurons or oligodendrocytes in a patient in need thereof, said method comprising introducing the cell population of embodiment 1 into said patient, wherein said cells are capable of differentiating into nervous tissue.

40. The method of embodiment 39, wherein said population is introduced to prevent neural degeneration.

41. The method of embodiment 39, wherein said population is introduced to ameliorate neural damage or degeneration.

42. The method of embodiment 39, wherein greater than about 10% of the cells within said population express a cell surface marker selected from the group

consisting of CD34, CD38, CD41, CD45, CD117, HLA-DR, and combinations thereof, wherein said cells are capable of multilineage development.

43. The method of embodiment 39, wherein said population does not substantially express cell surface markers selected from the group consisting of CD3, CD7, CD10, CD13, CD14, CD19, CD33, CD35, CD56, CD90, CD127, CD133, CD 135, CD138, glycophorin A, and combinations thereof.

44. The method of embodiment 39, wherein greater than about 10% of the cells within said population are side scatter channel low (SSC^{lo}).

45. The method of embodiment 39, wherein greater than about 10% of the cells within said population are side scatter channel high (SSC^{hi}).

46. A method of producing cardiomyocytes in a patient in need thereof, said method comprising introducing the cell population of embodiment 1 into said patient, wherein said cells are capable of differentiating into heart tissue.

47. The method of embodiment 46, wherein said population is introduced to prevent ischemic heart injury.

48. The method of embodiment 46, wherein said population is introduced to ameliorate ischemic heart injury.

49. The method of embodiment 46, wherein greater than about 10% of the cells within said population express a cell surface marker selected from the group consisting of CD34, CD38, CD41, CD45, CD117, HLA-DR, and combinations thereof, wherein said cells are capable of multilineage development.

50. The method of embodiment 46, wherein said population does not substantially express cell surface markers selected from the group consisting of CD3,

CD7, CD10, CD13, CD14, CD19, CD33, CD35, CD56, CD90, CD127, CD133, CD 135, CD138, glycophorin A, and combinations thereof.

51. The method of 46, wherein greater than about 10% of the cells within said population are side scatter channel low (SSC^{lo}).

52. The method of 46, wherein greater than about 10% of the cells within said population are side scatter channel high (SSC^{hi}).

53. A cell population of bone-marrow-derived, ALDH^{br} stem cells, wherein said cells are capable of multilineage development.

54. The cell population of embodiment 53, wherein greater than about 10% of the cells within said population express a cell surface marker selected from the group consisting of CD34, CD38, CD41, CD45, CD105, CD117, HLA-DR, and combinations thereof, wherein said cells are capable of multilineage development.

55. The cell population of embodiment 54, wherein greater than about 10% of the cells within said population express at least CD105.

56. The cell population of embodiment 55, wherein greater than about 40% of the cells within said population express at least CD105.

57. The cell population of embodiment 53, wherein said population does not substantially express cell surface markers selected from the group consisting of CD3, CD7, CD10, CD13, CD14, CD19, CD33, CD35, CD56, CD90, CD127, CD133, CD 135, CD138, glycophorin A, and combinations thereof.

58. The cell population of embodiment 53, wherein greater than about 10% of the cells within the population are side scatter channel low (SSC^{lo}).

59. The cell population of embodiment 53, wherein greater than about 10% of the cells within the population are side scatter channel high (SSC^{hi}).

60. The cell population of embodiment 53, wherein said population is capable of engrafting a mammal.

61. The cell population of embodiment 61, wherein said population is capable of engrafting hematopoietic cells.

62. The cell population of embodiment 61, wherein said population is capable of engrafting a SCID/NOD mouse spleen with human B cell precursors.

63. The cell population of embodiment 60, wherein said population is capable of engrafting human thymus tissue transplanted into SCID/hu Thy mice with T cell precursors.

64. The cell population of embodiment 63, wherein said T cell precursors are capable of developing into T cells expressing CD4 or CD8.

65. The cell population of embodiment 60, wherein said population is capable of engrafting mesenchymal cells.

66. The cell population of embodiment 65, wherein said population is capable of engrafting tissue selected from the group consisting of bone marrow stroma, bone, cartilage, tendon, fat, smooth muscle, cardiac muscle, skeletal muscle, nerves, oligodendrocytes, fibroblasts, endothelium, and combinations thereof.

67. A composition comprising the cell population of embodiment 53 in a pharmacologically acceptable carrier.

68. A method of reconstituting blood tissue in a patient in need thereof, said method comprising introducing the cell population of embodiment 53 into said patient, wherein said cells are capable of engraftment.

69. The method of embodiment 68, wherein said patient is in need of treatment for bone marrow ablation.

70. The method of embodiment 68, wherein said patient is in need of treatment for cancer.

71. The method of embodiment 68, wherein said patient is in need of treatment for sequelae related to cancer therapy.

72. The method of embodiment 68, wherein greater than about 10% of the cells within said population express a cell surface marker selected from the group consisting of CD34, CD38, CD41, CD45, CD105, CD117, HLA-DR, and combinations thereof, wherein said cells are capable of multilineage development.

73. The method of embodiment 72, wherein greater than about 10% of the cells within said population express at least CD105.

74. The method of embodiment 72, wherein greater than about 40% of the cells within said population express at least CD105.

75. The method of embodiment 68, wherein said population does not substantially express cell surface markers selected from the group consisting of CD3, CD7, CD10, CD13, CD14, CD19, CD33, CD35, CD56, CD90, CD127, CD133, CD 135, CD138, glycophorin A, and combinations thereof.

76. The method of embodiment 68, wherein greater than about 10% of the cells within said population are side scatter channel low (SSC^{lo}).

77. The method of embodiment 68, wherein greater than about 10% of the cells within said population are side scatter channel high (SSC^{hi}).

78. A method of repairing or regenerating a mesenchymal tissue in a patient in need thereof, said method comprising introducing the cell population of embodiment 53 into said patient.

79. The method of embodiment 78, wherein said mesenchymal tissue is selected from the group consisting of bone, cartilage, fat, endothelium, muscle, and combinations thereof.

80. The method of embodiment 79, wherein said cell population promotes neovascularization.

81. The method of embodiment 78, wherein said population is introduced to correct a bone defect.

82. The method of embodiment 78, wherein said population is introduced to correct a cartilage defect.

83. The method of embodiment 78, wherein greater than about 10% of the cells within said population express a cell surface marker selected from the group consisting of CD34, CD38, CD41, CD45, CD105, CD117, HLA-DR, and combinations thereof, wherein said cells are capable of multilineage development.

84. The method of embodiment 83, wherein greater than about 10% of the cells within said population express at least CD105.

85. The method of embodiment 83, wherein greater than about 40% of the cells within said population express at least CD105.

86. The method of embodiment 78, wherein said population does not substantially express cell surface markers selected from the group consisting of CD3, CD7, CD10, CD13, CD14, CD19, CD33, CD35, CD56, CD90, CD127, CD133, CD 135, CD138, glycophorin A, and combinations thereof.

87. The method of embodiment 78, wherein greater than about 10% of the cells within said population are side scatter channel low (SSC^{lo}).

88. The method of embodiment 78, wherein greater than about 10% of the cells within said population are side scatter channel high (SSC^{hi}).

89. A method of inducing immunological tolerance in a patient in need thereof, said method comprising introducing the cell population of embodiment 53 into said patient, wherein said cells are capable of downregulating alloantigen recognition and response.

90. The method of embodiment 89, wherein said population is introduced to prevent graft versus host disease.

91. The method of embodiment 89, wherein said population is introduced to ameliorate graft versus host disease.

92. The method of embodiment 89, wherein greater than about 10% of the cells within said population express a cell surface marker selected from the group consisting of CD34, CD38, CD41, CD45, CD105, CD117, HLA-DR, and combinations thereof, wherein said cells are capable of multilineage development.

93. The method of embodiment 92, wherein greater than about 10% of the cells within said population express at least CD105.

94. The method of embodiment 92, wherein greater than about 40% of the cells within said population express at least CD105.

95. The method of embodiment 89, wherein said population does not substantially express cell surface markers selected from the group consisting of CD3, CD7, CD10, CD13, CD14, CD19, CD33, CD35, CD56, CD90, CD127, CD133, CD 135, CD138, glycophorin A, and combinations thereof.

96. The method of embodiment 89, wherein greater than about 10% of the cells within said population are side scatter channel low (SSC^{lo}).

97. The method of embodiment 89, wherein greater than about 10% of the cells within said population are side scatter high (SSC^{hi}).

98. A method of producing neurons or oligodendrocytes in a patient in need thereof, said method comprising introducing the cell population of embodiment 53 into said patient, wherein said cells are capable of differentiating into nervous tissue.

99. The method of embodiment 98, wherein said population is introduced to prevent neural degeneration.

100. The method of embodiment 98, wherein said population is introduced to ameliorate neural damage or degeneration.

101. The method of embodiment 98, wherein greater than about 10% of the cells within said population express a cell surface marker selected from the group consisting of CD34, CD38, CD41, CD45, CD105, CD117, HLA-DR, and combinations thereof, wherein said cells are capable of multilineage development.

102. The method of embodiment 101, wherein greater than about 10% of the cells within said population express at least CD105.

103. The method of embodiment 101, wherein greater than about 40% of the cells within said population express at least CD105.

104. The method of embodiment 98, wherein said population does not substantially express cell surface markers selected from the group consisting of CD3, CD7, CD10, CD13, CD14, CD19, CD33, CD35, CD56, CD90, CD127, CD133, CD 135, CD138, glycophorin A, and combinations thereof.

105. The method of embodiment 98, wherein greater than about 10% of the cells within said population are side scatter channel low (SSC^{lo}).

106. The method of embodiment 98, wherein greater than about 10% of the cells within said population are side scatter channel high (SSC^{hi}).

107. A method of producing cardiomyocytes in a patient in need thereof, said method comprising introducing the cell population of embodiment 53 into said patient, wherein said cells are capable of differentiating into heart tissue.

108. The method of embodiment 107, wherein said population is introduced to prevent ischemic heart injury.

109. The method of embodiment 107, wherein said population is introduced to ameliorate ischemic heart injury.

110. The method of embodiment 107, wherein greater than about 10% of the cells within said population express a cell surface marker selected from the group consisting of CD34, CD38, CD41, CD45, CD105, CD117, HLA-DR, and combinations thereof, wherein said cells are capable of multilineage development.

111. The method of embodiment 110, wherein greater than about 10% of the cells within said population express at least CD105.

112. The method of embodiment 110, wherein greater than about 40% of the cells within said population express at least CD105.

113. The method of embodiment 107, wherein said population does not substantially express cell surface markers selected from the group consisting of CD3, CD7, CD10, CD13, CD14, CD19, CD33, CD35, CD56, CD90, CD127, CD133, CD 135, CD138, glycophorin A, and combinations thereof.

114. The method of embodiment 107, wherein greater than about 10% of the cells within said population are side scatter channel low (SSC^{lo}).

115. The method of embodiment 107, wherein greater than about 10% of the cells within said population are side scatter channel high (SSC^{hi}).

116. A method of screening a compound for its ability to promote differentiation, growth, cytotoxicity, apoptosis, or engraftment of stem cells, comprising the steps of:

- a) isolating ALDH^{br} stem cells from a stem cell source;
- b) further selecting a subpopulation of cells expressing CD105; and
- c) contacting said subpopulation of cells with said compound.

117. A method of screening a compound for its ability to promote differentiation, growth, cytotoxicity, apoptosis, or engraftment of stem cells, comprising the steps of:

- a) isolating ALDH^{br} stem cells from bone marrow;
- b) further selecting a subpopulation of ALDH^{br} stem cells expressing markers selected from the group consisting of CD34, CD38, CD41, CD45, CD105, CD117, HLA-DR, and combinations thereof; and

c) contacting said subpopulation of ALDH^{br} cells with said compound.

118. A kit comprising a detectable ALDH substrate disposed in a container, and antibodies specific for cell surface markers selected from the group consisting of CD34, CD38, CD41, CD45, CD105, CD117, HLA-DR, and combinations thereof, disposed in a container.

119. The kit of embodiment 118, wherein the ALDH substrate is BODIPY aminoacetaldehyde diethyl acetal or BODIPY aminoacetaldehyde.

BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 shows the scatterplots from Example 1 for the isotype and fluorescent controls.

Figure 2 shows the expression of surface markers from Example 1.

Figure 3 shows a gated scatterplot from Example 1 of the expression of ALDH^{br}SSC^{lo} in different grafts.

Figure 4 shows the expression of ALDH^{br}SSC^{lo} cell populations in different grafts for each test subject from Example 1.

Figure 5 shows the scatterplots from Example 1 for ALDH^{br} and ALDH^{dim} cells from bone marrow, umbilical cord blood, and mobilized peripheral blood are shown for cells expressing CD34, CD45, CD105, and CD133.

Figure 6 shows a comparison of CD34⁺ALDH^{br} subpopulations from Example 1.

DETAILED DESCRIPTION OF THE INVENTION

The present invention is directed to populations of stem cells, methods for isolating these stem cell populations, and methods of using these stem cell populations to reconstitute, regenerate, or repair tissue. These stem cell populations comprise stem and progenitor cells (SPC) that are ALDH^{br}, and, thus, contain most or all of the stem cells present in a stem cell source. The ALDH^{br} cell population further comprises subpopulations of cells that express combinations of cell markers, such as CD105, or have low granularity (as measured by flow cytometry), that serve to identify those subpopulations useful for particular embodiments of the invention such as for reconstituting, regenerating, or repairing a disease of interest or for manufacturing kits.

Using a combination of cell surface markers and others markers such as intracellular enzymes and the light scattering properties of the cells, stem cell grafts of the invention can be advantageously “tailored” for particular therapeutic uses. Also, because some of these unique stem cell populations represent certain cell lineages, the populations can be used to both selectively reconstitute certain cell lineages *in vivo*. Stem cells that give rise to hematopoietic lineages can be used to increase the concentration and potency of stem cell grafts and, thereby, decrease toxicity. Advantageously, these cells can be sorted from autologous bone marrow and peripheral

blood, thus further reducing the chance of rejection and increasing the efficacy of stem cell grafts. Stem cells that give rise to mesenchymal tissues such as bone, nerves, oligodendrocytes, muscles, vasculature, bone marrow stroma, and dermis can be used to repair or replace diseased or damaged tissues. Thus, the novel combination of CD markers disclosed herein confers such additional advantages as identification of the stem cell sources that are functionally and quantitatively best for use in isolating cells for stem cell grafts.

By “cell surface marker” is intended a protein expressed on the surface of a cell, which is detectable via specific antibodies. Cell surface markers that are useful in the invention include, but are not limited to, the CD (clusters of differentiation) antigens CD1a, CD2, CD3, CD5, CD7, CD8, CD10, CD13, CD14, CD16, CD19, CD29, CD33, CD34, CD35, CD38, CD41, CD45, CD56, CD71, CD73, CD90, CD105, CD115, CD117, CD124, CD127, CD130, CD133, CD135, CD138, CD144, CD166, HLA-A, HLA-B, HLA-C, HLA-DR, and glycophorin A. By “intracellular marker” is intended expression of a gene or gene product such as an enzyme that is detectable. For example, aldehyde dehydrogenase (ALDH) is an intracellular enzyme that is expressed in most hematopoietic stem cells. It can be detected via flow cytometry by using fluorescent substrates. By side scatter channel (SSC) and forward scatter channel (FSC) is intended the light refractive and reflective properties of a cell, respectively, as measured by flow cytometry, i.e., as cell granularity increases, the side scatter increases and as cell diameter increases, the forward scatter increases. Thus, the embodiments of the invention include selection of stem cell populations based on combinations of cell surface markers, intracellular markers, and the reflective and refractive properties of cells obtained from a stem cell source.

The populations of stem cells disclosed herein can comprise ALDH^{br} cells that can be sorted based on the positive expression of cell surface markers such as CD29, the integrin β 1 subunit expressed on most cells; CD34, a highly glycosylated type I transmembrane protein expressed on 1-4% of bone marrow cells; CD38, a type II transmembrane protein found on immature T and B cells but not most mature peripheral lymphocytes; CD41, the integrin α IIb subunit that is expressed on platelets and megakaryocytes; CD45, the leukocyte common antigen found on all cells of

hematopoietic origin; CD73, an ecto-5'-nucleotidase differentially expressed on subsets of mature lymphocytes and some endothelial and epithelial cells; CD105, a disulfide-linked homodimer found on endothelial cells but absent from most T and B cells; CD115, the macrophage colony stimulating factor receptor expressed primarily on cells of the mononuclear-phagocytic lineage; CD117, the c-kit ligand receptor found on 1-4% of bone marrow stem cells; CD133, a pentaspan transmembrane glycoprotein expressed on primitive hematopoietic progenitor cells; CD135, the Flt3 ligand receptor that is expressed on CD34⁺ hematopoietic stem cells; CD138, an extracellular matrix receptor expressed on immature B cells and plasma cells; CD144, the VE-cadherin molecule that organizes adherens junction in endothelial cells; CD166, found on thymic epithelium, activated T cells, and neurons; HLA-A, HLA-B, HLA-C, forms of the MHC Class I molecule; HLA-DR, the MHC Class II molecule; the VEGF receptor, found on hematopoietic stem cells and vascular endothelium; and any combinations of these markers.

In one embodiment, the population of cells comprises ALDH^{br} cells wherein at least 10% to 100% of the cells express at least CD41, at least CD105, or at least HLA-DR. In another embodiment, the population of cells comprises ALDH^{br} SSC^{lo} cells wherein at least 10% to 100% of the cells express any combination of CD41, CD105, and HLA-DR.

In some embodiments, the lack of expression of a cell surface marker defines stem cell populations of the invention. Examples include populations of stem cells comprising ALDH^{br} SSC^{lo} cells negative for expression of the following markers: CD1a, a lymphoid marker structurally similar to MHC Class I; CD2, a pan lymphoid marker associated with antigen recognition; CD3, a member of the T cell receptor complex; CD5, expressed on mature T and B cells; CD7, an early T cell lineage marker; CD10, a type II membrane metalloprotease expressed on early T and B cell precursors; CD13, a type II membrane metalloprotease expressed on granulocytes monocytes and their precursors; CD14, a GPI-linked protein expressed mainly on myelomonocytic lineage cells; CD19, a component of the B cell antigen signaling complex; CD33, a sialic acid binding protein that is absent from pluripotent stem cells but appears on myelomonocytic precursors after CD34; CD35, the complement receptor type 1, which is expressed on

many lymphoid and myelomonocytic cells; CD56, an isoform of the neural adhesion molecule found exclusively on natural killer (NK) cells; CD90, a GPI-cell anchored molecule found on prothymocyte cells in humans; CD127, the high affinity interleukin 7 receptor expressed on lymphocytes; CD138, an extracellular matrix receptor found on immature B cells and plasma cells; glycophorin A, a sialoglycoprotein present on human red blood cells and embryoid precursors; and any combinations of these markers. In some embodiments stem cells are Lin⁻; these stem cells do not yet express lineage-commitment cell surface markers, and therefore comprise a greater number of hematopoietic stem cells. By the term "Lin⁻" is intended that the cell lacks expression of the cell surface markers CD2, CD3, CD14, CD16, CD19, CD24, CD56, CD66b, CD14, and glycophorin A (gly A).

Based on these unique cell surface marker signatures, individual stem cell populations having unique functional characteristics have been identified.

In some embodiments at least 10%, 20%, or 30% of the ALDH^{br} cells within a stem cell population of the invention express the cell markers of interest; in other embodiments at least 40%, 50%, or 60% of the ALDH^{br} cells within the stem cell population express the cell markers of interest; in yet other embodiments at least 70%, 80%, or 90% of the ALDH^{br} cells within the stem cell population express the cell markers of interest; in still other embodiments at least 95%, 96%, 97%, 98%, 99%, or even 100% of the ALDH^{br} cells within the stem cell population express the cell markers of interest. While the isolation of purified cell population from bone marrow is specifically exemplified herein, the isolation of such cells from other sources, including umbilical cord blood, peripheral blood, and fetal liver, is also contemplated.

Selective methods known in the art and described herein can be used to further characterize SPC, including ASPC. Commonly, sources of SPC and ASPC are reacted with monoclonal antibodies, and subpopulations of cells expressing cognate cell surface antigens are either positively or negatively selected with immunomagnetic beads by complement mediated lysis, agglutination methods, or fluorescence activated cell sorting (FACS). The functional attributes of the resulting subpopulations with a defined cell surface phenotype are then determined using a colony-forming assay. Once the

phenotype of cells that do and do not have SPC or ASPC activity is known, these methods can be used to select appropriate SPC or ASPC for therapeutic transplantation.

If desired, a large proportion of terminally differentiated cells may be removed by initially using a negative selection separation step. For example, magnetic bead separations may be used initially to remove large numbers of lineage committed cells. Desirably, at least about 80%, usually at least about 70% of the total cells, will be removed.

Procedures for cell separation may include, but are not limited to, positive or negative selection by means of magnetic separation using antibody-coated magnetic beads, affinity chromatography, cytotoxic agents joined to a monoclonal antibody or used in conjunction with a monoclonal antibody, including, but not limited to, complement and cytotoxins, and "panning" with antibody attached to a solid matrix, e.g., plate, elutriation, or any other convenient technique.

Techniques providing accurate cell separation include, but are not limited to, flow cytometry, which can have varying degrees of sophistication, e.g., a plurality of color channels, low angle and obtuse light scattering detecting channels, impedance channels, and the like. The antibodies for the various dedicated lineages may be illuminated by different fluorochromes. Fluorochromes that may find use in a multicolor analysis include phycobiliproteins, e.g., phycoerythrin and allophycocyanins; fluorescein; and Texas red. The cells may also be selected against dead cells, by employing dyes that selectively accumulate in dead cells (e.g., propidium iodide and 7-aminoactinomycin D (7-AAD)). Preferably, the cells are collected in a medium comprising about 2% fetal calf serum (FCS) or 0.2% bovine serum albumin (BSA). See, for example, Fallon *et al.* (2003) *Br. J. Haematol.* 121:1, herein incorporated by reference.

Other techniques for positive selection may be employed, which permit accurate separation, such as affinity columns, and the like. The method of choice should permit the removal of the non-progenitor cells to a residual amount of less than about 20%, less than about 15%, less than about 10%, or less than about 5% of the desired population of stem cells.

The stem cell populations of the invention can be isolated from stem cell sources using a variety of methods, including those described herein and exemplified below. For

example, in one embodiment, a two-step procedure can be used. For example in a first step, stem cells are positively selected by first sorting for cellular expression of ALDH^{br}, for example by using FACS. In a second step, cells are either positively selected by sorted for expression of cell surface markers such as CD105, HLA-DR, CD41, or negatively sorting for lack of expression of cell surface markers such as lineage commitment markers.

Alternatively, a three-step isolation procedure can be used. In a first step, mature mononuclear cells, for example, which express any of the different surface antigens associated with specific lineage commitment, can be eliminated using a commercially available kit from StemCell Technologies. The resulting preparation, designated Lin⁻, is essentially devoid of cells expressing any of the following surface antigens: glycoprotein A (glyA), CD2, CD3, CD14, CD16, CD19, CD24, CD56, and CD66b. In a second step, a subset of Lin⁻ cells can be sorted by flow cytometry after staining with, for example, BODIPY aminoacetaldehyde, to obtain ALDH^{br} Lin⁻ and ALDH^{lo} Lin⁻. In a third step, CD34⁺ cells can be sorted from the ALDH^{br} Lin⁻ fraction using, for example, flow cytometry, to yield ALDH^{br} CD34⁻ Lin⁻ cells and ALDH^{br} CD34⁺ Lin⁻ cells. This preparation can further be sorted based on expression of CD13, CD33, CD35, CD38, CD41, CD45, CD90, CD105, CD117, CD133, CD135, CD138, HLA-DR, and any combinations thereof.

In an alternative embodiment, the isolation procedure involves the use of a single negative depletion step. In accordance with this procedure, cells expressing Lin lineage markers are eliminated in a single step using appropriate antibodies.

Regardless of the isolation procedure used, the resulting cells have properties of multilineage stem cells. By “multilineage stem cells” is intended cells capable of multilineage development into cells such as mesenchymal stem cells (MSC), hematopoietic stem cells (HSC), or self-renewing progenitor cells that are themselves capable of MSC or HSC development. By “multilineage development” is intended capable of developing into any differentiated tissue including, but not limited to, blood cells (including lymphocytes, myelocytes, erythrocytes, and platelets); bone (including osteocytes, osteoblasts, and osteoclasts); marrow stroma; cartilage (including chondrocytes, chondroblasts, and chondroclasts); cardiac, smooth, and skeletal muscle

(including myocytes and cardiomyocytes); tendon; nerves (including oligodendrocytes and neurons); vascular tissue (including angiocytes and the endothelium); fat (including adipocytes); fibroblasts; liver cells; gut and lung epithelial cells; cornea (including corneocytes; and dermis (including dermal cells). By “hematopoietic stem cells” is intended stem cells that differentiate into blood cells. By “mesenchymal stem cells” is intended stem cells that differentiate into non-blood cells. SPC and ASPC comprise multilineage stem cells capable of developing into either mesenchymal stem cells or hematopoietic stem cells.

Hematopoietic stem cells do not express cell surface markers for immature and mature lymphocytes, erythrocytes, or myeloid cells, unlike lineage-committed precursors. More specifically, they have no detectable expression of surface markers of myeloid cells (CD33), B-lymphocytes (CD19, HLA-DR), T-lymphocytes (CD3, CD4, CD8), NK cells (CD16, CD56), or erythroid cells (CD71), where expression is detected using fluorescence-conjugated antibodies. Hematopoietic stem cells also do not have detectable expression of the early lymphoid markers CD10, CD1a, CD2, or CD5, which are present on committed lymphoid precursors. Advantageously, the primitive hematopoietic stem cells of the invention do not have detectable expression of any of the following antigens: CD3, CD4, CD8, CD13, CD14, CD16, CD19, CD25, CD33, CD56, CD71, CD124, CD130, CD135, CD138, or glyA. Preferably, the ALDH^{br} stem cells are also capable of sustaining long-term hematopoiesis in mice (see, for example, Goodell *et al.* (1996) *J. Exp. Med.* 183:1797). More preferably, they are in G₀, a cell cycle state that is believed to be a property of primitive hematopoietic cells (Spangrude *et al.* (1990) *Proc. Natl. Acad. Sci. USA* 87:7433).

In one embodiment, the stem cells of the invention comprise ALDH^{br} cells derived from bone marrow. These ALDH^{br} cells can be used to generate any cell of the hematopoietic lineage, including, but not limited to, myeloid cells (such as platelets, megakaryocytes, and red bloods cells) and lymphoid cells (such as T cells, B cells, NK cells, and antigen presenting cells). In an alternative embodiment, the stem cell of the invention can comprise ALDH^{br} CD105⁺ cells isolated from any stem cell source, not just bone marrow. These cells are also capable of generating any cell of the hematopoietic lineage, including, but not limited to, myeloid cells (such as platelets, megakaryocytes,

and red blood cells) and lymphoid cells (such as T cells, B cells, NK cells, and antigen presenting cells). In an alternative embodiment, the ALDH^{br} cells derived from bone marrow or the ALDH^{br} CD105⁺ cells isolated from any stem cell source comprise cells with a low granularity as measured in the side scatter channel of a flow cytometer (SSC^{lo}).

Mesenchymal stem cells (MSC) have been characterized using panels of antibodies much like hematopoietic stem cells. MSC generally lack expression of CD14, CD34, and CD45. MSC are generally positive for CD105 and CD73. Other markers used by researchers to identify cultured mesenchymal cells include positive expression of such markers as CD29, Thy-1 (CD90), CD115, CD144, CD166, and HLA-A,B, or C. Functionally, MSC can be tested *in vitro* for their ability to differentiate into adipogenic, osteogenic, myogenic, and chondrogenic cell colonies.

As disclosed *supra*, the stem cells of the invention comprise ALDH^{br} cells derived from bone marrow. These ALDH^{br} cells can be also used to at least generate any cell of the mesenchymal lineage, including, but not limited to, bone (including osteocytes, osteoblasts, and osteoclasts), marrow stroma, cartilage (including chondrocytes, chondroblasts, and chondroclasts), cardiac, smooth, and skeletal muscle (including myocytes and cardiomyocytes), tendon, nerves (including oligodendrocytes and neurons), vascular tissue (including the endothelium), fat (including adipocytes), fibroblasts, and dermis. In an alternative embodiment, the stem cell of the invention can comprise ALDH^{br} CD105⁺ cells isolated from any stem cell source, not just bone marrow. These cells are also capable of generating any cell of the mesenchymal lineage, including, but not limited to, bone (including osteocytes, osteoblasts and osteoclasts), marrow stroma, cartilage (including chondrocytes, chondroblasts, and chondroclasts), cardiac, smooth, and skeletal muscle (including myocytes and cardiomyocytes), tendon, nerves (including oligodendrocytes and neurons), vascular tissue (including the endothelium), fat (including adipocytes), fibroblasts, and dermis. In an alternative embodiment the ALDH^{br} cells derived from bone marrow or the ALDH^{br} CD105⁺ cells isolated from any stem cell source comprise cells with a low granularity as measured in the side scatter channel of a flow cytometer (SSC^{lo}).

Functional assays for stem cells include both *in vitro* and *in vivo* methods. For *in vitro* tests, cultures are seeded with cells from tissues under conditions that favor the differentiation of SPC or ASPC into specific types of tissue cells. See, for example, Eaves, "Assays of Hematopoietic Progenitor Cells" in Williams (1995) *Hematology* at L22-6 (5th ed., E. Beutler *et al.* eds.); Petzer *et al.* (1996) *Proc. Natl. Acad. Sci. USA* 93:1470; Mayani *et al.* (1993) *Blood* 81:3252; Hogge *et al.* (1997) *Br. J. Haematol.* 96:790; Lazarus *et al.* (1995) *Bone Marrow Transplant.* 16:557; Pittenger *et al.* (1999) *Science* 284:143.

Usually, ASPC will multiply a few times in culture as they differentiate to form a colony of differentiated cells. In some assays, such as assays involving blood cell differentiation, the developmental potential of a cell that initiated the colony is inferred from the range of phenotypes within the colony. Alternatively, cell cultures may be sampled and subcultured into a different culture system or into animal models, such as immunodeficient NOD-SCID mice or sheep embryos, to determine if additional phenotypes can be induced from APSC that remain in the individual colony. See, for example, Almeida-Porada *et al.* (2000) *Blood* 95:3620; Lewis *et al.* (2001) *Blood* 97:3441; Rice *et al.* (2000) *Transplantation* 69:927; Ishikawa *et al.* (2002) *Exp. Hematol.* 30:488. ASPC that give rise to multiple phenotypes are usually considered more primitive, multipotent progenitors than cells that give rise to fewer phenotypes.

Based on these assays, the choice of a starting stem cell source coupled with the specific lineage-preferred cell population allows for a variety of therapeutic applications using one or a few ASPC types. For example, ALDH^{br} cell populations that have been isolated by sorting BM include more mesenchymal stem cells capable of giving rise to non-hematopoietic tissues than do ALDH^{br} cell populations derived from MPB or UCB based on the increased percentage of cells expressing early mesenchymal stem cell markers such as CD105. See, Example 1, Table 1, herein below.

The stem cell populations of the present invention have application in a variety of therapies and diagnostic regimens. They are preferably diluted in a suitable carrier such as buffered saline before injection. The cells may be administered in any physiologically acceptable vehicle, normally intravascularly, although they may also be introduced into bone or other convenient site where the cells may find an appropriate site for regeneration

and differentiation (e.g., thymus). Usually, at least 1×10^5 cells/kg and preferably 1×10^6 cells/kg or more will be administered. See, for example, Sezer *et al.* (2000) *J. Clin. Oncol.* 18:3319 and Siena *et al.* (2000) *J. Clin. Oncol.* 18:1360. The cells may be introduced by injection, catheter, or the like. If desired, additional drugs such as 5-fluorouracil and/or growth factors may also be co-introduced. Suitable growth factors include, but are not limited to, cytokines such as IL-2, IL-3, IL-6, IL-11, G-CSF, M-CSF, GM-CSF, gamma-interferon, and erythropoietin. In some embodiments, the cell populations of the invention can be administered in combination with other cell populations that support or enhance engraftment, by any means including but not limited to secretion of beneficial cytokines and/or presentation of cell surface proteins that are capable of delivering signals that induce stem cell growth, homing, or differentiation. In these embodiments, less than 100% of the graft population comprise the ALDH^{br} stem cells.

As the isolated cells of the invention are capable of engrafting hematopoietic stem cells, they are suitable for both transplantation and gene therapy purposes. Since the populations of cells can be sorted into both lineage-committed and lineage-uncommitted cells, each population will have different, but non-mutually exclusive, therapeutic applications. In some embodiments, lineage-committed cells will be useful for reconstitution or augmentation of lymphoid and/or myeloid cell populations. For example, indications to be treated include immunodeficiencies and stem cell transplantations. The progenitor cells are particularly useful during stem cell transplantation to decrease the lag time between the transplantation and repopulation of the hematopoietic cells. Methods of obtaining the progenitor cells are described herein. Methods of administering stem cells to patients are within the skill of one in the art as noted herein above.

In some embodiments, identification of a lymphoid-restricted progenitor cell population allows assessment of this population as to the origin of disease, e.g., leukemia, and therefore is useful in purging such cells from an autologous graft. In other embodiments, lineage-committed precursor cells may be useful as a cellular graft to increase the patient's ability to mount an immune response. In some instances, it may be advantageous to use precursors in a vaccine strategy whereby the cells are loaded with

antigen and injected to induce a specific immune response. Alternatively, the injected cells may be used to induce tolerance to a specific exogenously introduced antigen.

Lineage-committed precursors may be used for a variety of gene therapy approaches where expression of the exogenous genetic capability is desired in lymphoid and/or myeloid lineages. Lineage-committed progenitor cells will also be preferred in cases where it is desired to have temporary rather than permanent expression of the exogenous genetic capability. Also, in some instances, gene transfer is likely to be more efficient in lineage-committed progenitors because progenitor cells cycle more actively than stem cells and retroviral vectors are known to require actively cycling cells for efficient integration of the recombinant DNA.

In addition, it would be advantageous to use lymphoid progenitors compared to using mature lymphoid cells for gene therapy. Currently, T cell gene therapy requires *ex vivo* expansion of T cells with cytokines. Upon reinfusion, the modified T cells often do not home properly to their target organs and may become trapped in (and cleared by) the lungs, liver, or spleen. This improper homing may be due to alteration of the membranes during the *ex vivo* processing, downregulation of homing receptors, or the like. Use of modified progenitor cells would obviate the necessity of *ex vivo* expansion of the effector T cells, and thus obviate concerns of altered trafficking and persistence *in vivo*. In addition, the use of modified progenitors will allow amplification in progeny cell numbers, thereby reducing the need for *ex vivo* expansion and reducing the frequency of administration.

In other embodiments, it is preferable to use non-lineage-committed cells or more primitive stem cells. Non-lineage committed cells are useful for treating diseases (and related therapeutic sequelae) caused by the impaired function of bone marrow cells. Examples of such diseases include, but are not limited to, lymphoma, multiple myeloma, breast cancer, testicular cancer, leukemias, congenial hemolytic anemias (e.g., thalassanemia), and some immunodeficiency diseases, e.g., acute leukemia, Hodgkin's and non-Hodgkin's lymphoma, and neuroblastoma. It has also been shown that the hematologic toxicity sequelae observed with multiple cycles of high-dose chemotherapy is relieved by conjunctive administration of autologous hematopoietic stem cells. Diseases for which reinfusion of stem cells has been described include acute leukemia,

Hodgkin's and non-Hodgkin's lymphoma, neuroblastoma, testicular cancer, breast cancer, multiple myeloma, thalassemia, and sickle cell anemia (Cheson *et al.* (1989) *Ann. Intern. Med.* 30 110:51; Wheeler *et al.* (1990) *J. Clin. Oncol.* 8:648; Takvorian *et al.* (1987) *N. Engl. J. Med.* 316:1499; Yeager, *et al.* (1986) *N. Eng. J. Med.* 315:141; Biron *et al.* (1985) In *Autologous Bone Marrow Transplantation: Proceedings of the First International Symposium*, Dicke *et al.*, eds., p. 203; Peters (1985) *ABMT*, *id.* at p. 189; Barlogie, (1993) *Leukemia* 7:1095; Sullivan, (1993) *Leukemia* 7:1098-1099).

For example, in cancer patients, sorting stem cells of the invention such as ALDH^{br} bone marrow derived cells or ALDH^{br} CD105⁺ cells from any source separates stem cells from cancer cells prior to reintroduction into the patient. In these patients undergoing autologous transplantation, such separation can be used to reduce the chance that cancer cells are returned to the patient (Jones *et al.* (1987) *Blood* 70:1490; Colvin In: *Hematopoietic Cell Transplantation* 217 (Forman ed., 1999), and Russo and Hilton (1989) in *Enzymology and Molecular Biology of Carbonyl Metabolism* at 65 (Weiner and Flynn, eds.). In another embodiment, purified autologous ALDH^{br} stem cells of the invention can be *ex vivo* expanded prior to reintroduction into the patient to hasten lymphoid, erythroid, and platelet engraftment. *Ex vivo* expansion can be effected by growth in defined cytokines, on stromal layers, and/or in bioreactors (Emerson *et al.* (1996) *Blood* 87:3082). In addition, the incidence of graft failure can be reduced. This is beneficial for cancer patients undergoing autologous transplantation, for gene therapy, and for patients suffering from auto-immune disorders.

In some embodiments, the population of non-lineage committed stem cells of the invention may be used to hasten healing of injuries, repair injuries, and regenerate tissues. As the cell populations of the invention can differentiate into mesenchymal tissue, the stem cells of the invention can be used beneficially in a wide variety of diseases not associated with hematopoiesis, for example joint regeneration for arthritis; bone regeneration for osteoporosis, osteoimperfecta, and periodontitis; angiogenesis and endothelial cell regeneration for ischemic injury; muscle re-growth for degenerative muscle disease such as muscular dystrophy and cardiac ischemia; neural cell regrowth in spinal cord and brain injury; regeneration of oligodendrocytes in multiple sclerosis; as well as surgeries where tissue regrowth is beneficial such as plastic surgery, orthopedic

surgery, and surgical tissue removal. Thus, the mesenchymal stem cells of the invention can be used to produce, in some embodiments, new endothelium, new cardiomyocytes, new neurons and oligodendrocytes, and new bone and cartilage. For example, it has been shown that mesenchymal stem cells grow endothelium, and that mesenchymal grafts treat restenosis and ischemic heart injury. See, for example, Hristov *et al.* (2003) *Trends Cardiovasc Med.* 13:201; Sata (2003) *Trends in Cardiovasc. Med.* 13:249; Isner *et al.* (2001) *Ann. NY Acad. Sci.* 953:75; Abbott and Giordano (2003) *J. Nucl. Cardiol.* 10:403; Forrester *et al.* (2003) *Circulation* 108:1139. It has also been shown that mesenchymal stem cell grafts promote regrowth of damaged neurons after neurological damage or disease. See, for example, Gage (2000) *Science* 287:1433; Shin *et al.* (2001) *Blood* 98:2412; Zhao *et al.* (2003) *Brain Res. Protocols* 11:38. It has also been shown that mesenchymal stem cell grafts promote new bone and cartilage growth. See, for example, Pittenger *et al.* (1999) *Science* 284:143, and U.S. Patent No. 6,541,024.

In some embodiments, the isolated stem cell populations of the invention are suitable for use in gene therapy. For example, following isolation of autologous ALDH^{br} cells from bone marrow or ALDH^{br}CD105⁺ from any stem cell source, the isolated cells are exposed to a gene delivery vector, and the genetically modified stem cells are reinfused into the patient. Gene therapy protocols using genetically modified stem cells are known in the art. See, for example, Smith (1992) *Hematother.* 1:155. This approach can involve *ex vivo* culture or the use of vectors capable of transferring genes into non-dividing cells, thereby rendering *ex vivo* culture unnecessary. By the term "*ex vivo* gene therapy" is meant the *in vitro* transfection or retroviral infection of stem cells prior to introducing the transfected stem cells into a mammal. Gene therapy can be useful in treating, for example, congenital diseases, which include, but are not limited to, adenosine deaminase deficiency, recombinae deficiency, recombinae regulatory gene deficiency, and sickle cell anemia. In treating sickle cell anemia, for example, the mutant β -globin gene is replaced or supplemented with either the wild-type globin gene or an anti-sickling globin gene. In the treatment of cancer, drug resistance genes can be introduced into cells to confer resistance to cytotoxic drugs. This can reduce the incidence and severity of myelosuppression. For the treatment of infectious diseases, including HIV, anti-viral genes can be introduced into stem cell populations of the

invention so that they are rendered resistant to the virus (see, for example, Gilboa and Smith (1994) *Trends in Genetics* 10:139).

In some embodiments, isolation of ALDH^{br} cells from bone marrow or ALDH^{br}CD105⁺ from any stem cell source results in the elimination of pre-T-cells that cause graft versus host disease (GvHD). This elimination from the stem cell population of the invention can be expected to reduce the incidence and severity of GvHD in recipients of allogeneic transplants. See, for example, Ho and Soiffer (2001) *Blood* 98:3192.

Isolated ASPC can be *ex vivo* expanded to hasten neutrophil, erythroid, and platelet engraftment after allogeneic transplantation. In addition, the incidence of graft failure can be reduced. This is likely to be particularly important for recipients of umbilical cord blood transplants, where small cell doses limit the success of transplantation. Techniques for *ex vivo* expansion are well described in the art. See, for example, McNiece and Briddle (2001) *Exp. Hematol.* 29:3; McNiece *et al.* (2000) *Exp. Hematol.* 28:1186; Jaroscak *et al.* (2003) *Blood* 101:5061.

In some embodiments, successful engraftment with ALDH^{br} cells from bone marrow or ALDH^{br}CD105⁺ from any stem cell source can also be expected to induce tolerance to alloantigens. Prior induction of tolerance enhances the chance of permanent engraftment of subsequent solid organ transplants that originate from the same donor.

It will be appreciated that the stem cell populations of the present invention can be used as sources of new genes (e.g., for cytokines and cytokine receptors), including genes important in growth and development.

In addition to their application in treatment and diagnosis strategies, the stem cell populations of the present invention can be used in screening protocols to identify agents that can be used, for example, to promote differentiation or growth and/or engraftment of stem cells. In one such protocol, ALDH^{br} cells from bone marrow or ALDH^{br}CD105⁺ from any stem cell source are contacted with a test compound suspected of inducing differentiation, and the ability of the test compound to effect differentiation is determined (using, for example, microscopic and flow cytometric examination). In another screening protocol, ALDH^{br} cells from bone marrow or ALDH^{br}CD105⁺ from any stem cell source are contacted with a test compound suspected of inducing proliferation and/or

engraftment and the ability of the test compound to effect proliferation and/or engraftment is determined, for example, using *in vitro* long-term colony assays or *in vivo* immunodeficient mice models (e.g., SCID/NOD mice). See, for example, Peault *et al.* (1993) *Leukemia* 7:s98-101.

The invention also relates to kits that can be used to prepare the stem cell populations of the present invention, where, in some embodiments, the populations comprise ALDH^{br} cells from bone marrow or ALDH^{br}CD105⁺ from any stem cell source. The kits can comprise an ALDH substrate disposed in a single container and antibodies that can be used to effect direct isolation of the cells in separate containers. In a preferred embodiment, the kit includes at least antibodies specific for lineage-specific markers disposed within one or more container means. These antibodies can be used in conjunction with those present in the StemCell Technologies kit to achieve, for example, about 80% purification or higher. Advantageously, the kit also includes antibodies specific for HLA-DR. Any or all of antibodies specific for CD2, CD3, CD8, CD10, CD13, CD14, CD16, CD19, CD33, CD34, CD35, CD38, CD41, CD45, CD56, CD71, CD90, CD105, CD117, CD124, CD127, CD130, CD133, CD135, CD138, and glyA can also be included, disposed within at least one container means. The kit can also include, disposed within a container means, anti-CD7 antibodies.

The antibodies of the kits are disposed within a container means and the kit can further include ancillary reagents (e.g., buffers and the like) suitable for carrying out the isolation protocols.

The following examples are offered by way of illustration and not by way of limitation.

EXPERIMENTAL

Example 1: Tissue Processing and Phenotyping

Directly conjugated fluorescent antibodies were employed for analyses of cell surface antigens. Those used included antibodies directed against CD3 (Leu4; Becton Dickinson clone SK7); CD7 (Leu9; Becton Dickinson clone M-T701); CD10 (Beckman

Coulter/Immunotech clone ALB1); CD13 (Beckman Coulter/Immunotech clone SJ1D1); CD14 (Beckman Coulter/Immunotech clone RM052); CD19 (Leu12; Pharmingen clone WM15); CD33 (LeuM9; Becton Dickinson clone M-φP9); CD34 (Becton Dickinson clone 8G12); CD35 (Pharmingen clone E11); CD38 (Leu17; Becton Dickinson clone HB7); CD41 (Caltag clone VIPL3); CD45 (Miltenyi clone 5B1); CD56 (Leu19; Becton Dickinson clone My31); CD90 (Pharmingen; clone 5E10); CD105 (Caltag clone SN6); CD117 (Becton Dickinson clone 104D2); CD127 (Beckman Coulter/ Immunotech clone R34.34); CD133 (Miltenyi clone AC 133); CD135 (Pharmingen clone 4G8); CD 138 (IQ products clone B-B4); HLA-DR (Becton Dickinson clone L243); and glyA (Becton Dickinson clone GAR-2).

Fresh bone marrow samples from ten normal donors (Cambrex) were diluted in ammonium chloride 1:40 by volume (0.17 M NH_4Cl containing 10 mM Tris-HCl, pH 7.2 and 200 mM EDTA final concentration) and incubated at 4°C for 30 minutes to lyse erythrocytes. Cells were pelleted by centrifugation, resuspended at $1 \times 10^7/\text{ml}$, and reacted for aldehyde dehydrogenase activity using the ALDEFLUOR™ kit (StemCo Biomedical, Inc.). Cells were suspended in assay buffer, and 50 ml of substrate/ml were added to the cells and mixed immediately. 100 ml of cells were removed from the substrate tube and added to the diethylaminobenzaldehyde (DEAB) control tube. Cells were incubated at 37°C for 30 minutes. $3\text{-}4 \times 10^6$ cells were placed in each immunophenotype tube and incubated for an additional 15 minutes at 4°C with the appropriate monoclonal antibodies. After incubation, 1 ml of assay buffer was added to each tube to wash the cells, and the cells were pelleted by centrifugation. The supernatant was removed, and the cells were resuspended in 300-500 ml of assay buffer.

Fresh UBC samples were red blood cell depleted using hetastarch and ammonium chloride lysis or hetastarch and ficoll. 20 ml of cord blood was diluted with an equal volume of PBS and 8 mls of hetastarch (Heta Sep; Stem Cell Technologies, Inc.). Samples were mixed and incubated for 60 minutes at room temperature. The white blood cell fraction was removed from each tube and pelleted by centrifugation. Recovered cells were resuspended in 30 ml of ammonium chloride and incubated at 37°C for 15-30 minutes or passed over a ficoll gradient to remove contaminating red blood cells. Cells

were washed, diluted, and phenotyped with monoclonal antibodies and ALDEFLUOR™ as described above.

Fresh peripheral blood samples from normal donors or fresh mobilized peripheral blood (i.e., post G-CSF infusion) from patients were diluted 1:20 in ammonium chloride and incubated for 15-30 minutes at 37°C. See, Fields *et al.* (1994) *Cancer Control* 1:213 and Elfenbein *et al.* (1995) *Annal. NY Acad. Sci.* 770:315 for patient procedures. The recovered cells were then washed, diluted, and phenotyped with monoclonal antibodies and ALDEFLUOR™ as described above.

All human samples were obtained using technical and informed consent procedures that complied with relevant regulations and were approved by local institutional review boards.

Multiparameter phenotyping data was acquired on a Becton Dickinson FACSCalibur using an argon laser emission at 488 nm to excite the ALDEFLUOR™ reaction product (fluorescence detected in FL1 channel), phycoerythrin-conjugated antibodies to a large panel of markers (see Table 1; fluorescence detected in FL2 channel); 7-aminoactinomycin D was used to stain and discriminate dead cells (fluorescence detected in FL3 channel), and a HeNe laser was used to excite allophycocyanine-conjugated CD34 (fluorescence detected in FL4). Fluorescence was carefully compensated in multiparameter analysis to correct for spillover of signals among channels (see Table 1 for compensation controls). The ALDH^{br}SSC^{lo} cell population was defined by gating on forward and side scatter and FL1. A large data file was acquired such that >2000 events were contained in the ALDH^{br}SSC^{lo} gate.

Table 1 shows the percentage of the cells within the ALDH^{br}SSC^{lo} window that express the given marker and also the percentage of the total cells in the sample that express the marker that fall outside the ALDH^{br} window. Thus, CD7, primarily a T cell marker, and CD45, a pan-blood cell marker, fall primarily outside the window, while the percentage of stem cell markers such as CD34 and CD133 are increased within the window relative to ungated cells. Notably, not all of the cells expressing stem cell markers are in the window. Therefore, ALDH defines subsets of these stem cell markers, and the populations are different in the three sources. The distribution of the markers in the ALDH^{br} window varies both among tissues and among samples of a given tissue

(especially for cord blood), which suggests that the ALDH marker gives more information about the physiological and functional state of the stem cells.

Table 2 shows the percentage of CD34⁺ cells that also express each of the other markers shown in each tissue. In this table, all markers were analyzed with regard to CD34 and ALDH expression.

Table 3 shows the overall expression of the markers in the three tissues.

Table 4A-C summarizes the expression of CD markers in different gates for bone marrow, umbilical cord blood, and peripheral blood, respectively. The overall expression in the tissue, the expression in the ALDH^{br} gate, and the expression in the CD34⁺ gate are shown.

Within this gate (ALDH^{br}SSC^{lo}), it was observed that BM includes more cells capable of giving rise to non-hematopoietic tissues (e.g., CD34, CD45, CD105, CD117, and CD133) than populations derived from MPB or UCB based on the phenotypic profile. Moreover, CD105 is expressed to a greater extent in the bone marrow ALDH^{br} cell population than in other grafts. The CD105⁺ and CD133⁺ cells are very bright for ALDH in the bone marrow, even brighter than CD34. It is possible that CD133⁺ cells are CD105 positive, but that not all CD105⁺ cells are CD133 positive. CD105⁺ are backgated against CD133 to confirm this conclusion. Also, the SSC^{lo} window does not capture all of the CD105/CD133 ALDH^{br} cells in marrow, although it does capture the CD34⁺ cells, which is different in UCB and PBSC; CD105⁺ cells differentiate into ALDH^{dim} and a small population of ALDH very bright cells, while CD133 co-expresses with CD34 to a large extent in the SSC^{lo} window. The scatterplots and comparative data supporting these conclusions are shown in Figures 1-6

Mean and Standard Deviation Summary												
	Bone Marrow (n=10)				Cord Blood (n=7)				Peripheral Blood (n=16)			
	% of ALDH ^{br} cells expressing marker		% of cells expressing marker that are NOT ALDH ^{br}		% of ALDHbr cells expressing marker		% of cells expressing marker that are NOT ALDH ^{br}		% of ALDHbr cells expressing marker that are NOT ALDH ^{br}			
CD Marker	Mean	S.D.	Mean	S.D.	Mean	S.D.	Mean	S.D.	Mean	S.D.	Mean	S.D.
CD3	1.36	0.788	99.91	0.098	1.95	1.249	99.96	0.043	1.47	1.426	99.91	0.115
CD7	1.31	0.798	99.78	0.384	7.36	14.051	99.92	0.083	1.02	0.845	99.84	0.236
CD10	1.29	1.234	98.35	4.383	0.46	0.410	98.39	2.425	0.47	0.721	94.22	14.119
CD13	7.96	8.649	99.59	0.401	16.33	21.495	99.21	1.537	24.84	26.139	98.67	2.962
CD14	1.68	1.755	99.78	0.202	1.13	1.359	99.97	0.021	0.66	0.798	99.96	0.051
CD19	1.53	1.697	99.07	1.934	0.55	0.660	99.76	0.555	0.60	0.762	98.53	3.092
CD33	7.55	7.510	98.57	3.470	12.33	13.595	99.39	1.091	18.71	22.23	99.22	1.134
CD34	68.96	14.630	54.19	12.978	81.41	5.252	25.60	11.454	94.85	4.405	24.39	25.683
CD35	1.57	1.239	99.43	1.500	1.01	0.974	99.95	0.053	2.37	2.65	99.73	0.667
CD38	68.75	16.872	96.09	1.754	49.20	21.900	99.31	0.793	69.17	26.779	97.76	2.452
CD41-C	27.22	18.522	98.95	0.695	16.89	19.567	99.43	0.521	27.26	23.166	98.52	2.910
CD45-M	54.62	18.533	99.26	0.395	78.28	20.454	99.16	0.726	86.08	21.323	98.56	2.128
CD56-BD	2.16	1.398	97.39	2.762	4.64	9.899	99.40	0.936	1.62	2.269	98.55	3.868
CD90	3.22	3.532	93.87	6.810	8.00	14.188	98.88	0.984	2.52	2.356	89.10	15.818
CD105	10.28	11.173	75.18	22.390	0.82	0.686	94.87	10.439	1.03	1.188	93.06	16.374
CD117	43.78	23.037	72.54	14.873	45.96	26.423	43.51	8.297	18.99	18.045	45.61	17.208
CD127	0.74	0.688	76.77	19.260	0.39	0.255	88.66	14.759	0.54	0.657	73.89	27.193
CD133	4.06	4.626	61.29	32.003	29.61	21.858	37.21	14.145	36.80	26.149	26.23	16.888
CD135	0.84	0.704	84.93	18.976	1.09	1.428	90.93	15.926	0.58	0.717	81.56	19.431
CD138*	0.71	0.740	88.36	17.536	0.11		99.67		0.57	0.582	76.98	24.293
HLA-DR	38.43	16.437	97.46	1.620	65.79	20.136	98.02	1.952	64.89	26.688	95.61	4.872
GLY-A	5.67	4.942	98.02	1.942	1.46	0.479	98.47	1.994	0.69	0.694	96.06	6.321
ALDH	0.97	0.598			0.80	0.556			1.11	1.084		

*For Bone Marrow CD138: n=7

*For Cord Blood CD138: n=1

*For Peripheral Blood CD138: n=10

CD Marker	Bone Marrow (n=10)		Cord Blood (n=7)		Peripheral Blood (n=16)	
	Mean	STDEV	Mean	STDEV	Mean	STDEV
CD3	0.69	0.348	3.15	1.863	1.86	2.371
CD7	0.78	0.444	6.30	6.090	1.29	0.976
CD10	8.49	8.179	1.07	1.334	0.61	0.737
CD13	6.09	7.420	14.60	21.476	25.16	25.962
CD14	0.26	0.317	0.25	0.188	0.57	0.556
CD19	3.17	3.328	1.53	0.872	0.67	0.837
CD33	6.52	4.710	9.70	12.607	18.25	22.459
ALDH	47.19	11.882	74.40	11.454	83.41	11.946
CD35	0.48	0.424	1.22	1.078	2.06	2.040
CD38	69.92	20.245	47.80	22.845	67.65	27.161
CD41-C	25.35	19.138	16.51	21.023	26.94	23.251
CD45-M	51.03	22.604	75.61	25.070	84.27	23.342
CD56-BD	0.79	0.658	3.17	7.043	1.51	2.186
CD90	1.75	3.218	6.14	11.781	1.75	1.868
CD105	1.34	1.256	0.23	0.187	0.78	1.128
CD117	32.91	14.833	45.36	25.923	17.93	18.403
CD127	0.27	0.181	0.14	0.159	0.40	0.489
CD133	2.78	4.016	30.74	24.161	35.63	26.734
CD135	0.16	0.094	0.34	0.401	0.43	0.632
CD138	0.13	0.151	0.09	n/a	0.43	0.584
HLA-DR	49.76	18.578	64.26	22.024	62.44	26.427
GLY-A	0.38	0.228	0.85	0.556	0.51	0.611
ALDH	0.97	0.599	0.80	0.556	1.11	1.085
R3 iso	0.45	0.304	0.47	0.146	0.30	0.325
R5 iso	0.19	0.288	0.78	0.780	0.19	0.261
R2 pe iso	3.59	8.143	0.93	0.510	0.79	0.815
G7 pe iso	13.51	21.887	3.37	3.064	4.57	6.799

Table 2: Percentage of CD34+ Cells Expressing CD Marker Shown

CD Marker	Bone Marrow (n=10)		Cord Blood (n=6)		Peripheral Blood (n=16)	
	Mean	STDEV	Mean	STDEV	Mean	STDEV
CD3	14.72	3.727	43.69	14.643	21.81	18.080
CD7	8.38	2.719	43.76	17.460	14.85	15.860
CD10	3.73	2.290	0.79	0.847	0.72	0.933
CD13	19.42	8.313	22.12	10.275	45.08	27.214
CD14	5.24	1.362	22.62	3.440	25.43	11.399
CD19	2.68	1.751	6.45	3.990	4.69	8.967
CD33	15.51	9.257	23.73	3.985	25.98	17.523
CD34	1.40	0.500	0.90	0.517	0.98	1.082
CD35	9.58	8.285	20.08	8.910	41.67	21.568
CD38	17.66	3.871	71.88	18.735	36.31	18.805
CD41-C	26.32	14.980	22.33	5.725	41.81	19.781
CD45-M	70.51	13.339	88.84	12.651	96.93	4.692
CD56-BD	1.07	0.854	5.90	7.961	3.98	4.527
CD90	1.88	3.681	6.24	10.195	0.72	0.347
CD105	0.53	0.367	1.44	2.131	2.36	4.736
CD117	1.52	0.582	0.67	0.500	0.24	0.308
CD127	0.07	0.053	0.59	1.012	1.03	2.686
CD133	0.12	0.113	0.32	0.255	0.27	0.151
CD135	0.15	0.119	0.34	0.463	0.10	0.080
CD138	0.07	0.060	0.37	#DIV/0!	0.04	0.012
HLA-DR	14.92	2.690	31.27	6.338	21.49	10.799
GLY-A	9.16	5.722	3.07	2.110	0.47	0.379
ALDH	0.97	0.599	0.80	0.556	1.11	1.085
					n =9	
R3 iso	0.45	0.304	0.47	0.146	0.30	0.325
R5 iso	0.19	0.288	0.78	0.780	0.19	0.261
R2 pe iso	3.59	8.143	0.93	0.510	0.79	0.815
G7 pe iso	13.51	21.887	3.37	3.064	4.57	6.799
CD45 APC	79.57	11.861	89.39	10.992	96.96	2.178
AB45 APC	71.45	25.804	66.33	24.573	79.89	21.902

Table 3: Percentage of Ungated Cells Expressing Shown Markers

Bone Marrow						
CD Marker	ALDH CD+		CD34CD+		CD+	
	Mean %	St. Dev.	Mean %	St. Dev.	Mean %	St. Dev.
CD3	1.36	0.788	0.69	0.348	14.72	3.727
CD7	1.31	0.798	0.78	0.444	8.38	2.719
CD10	1.29	1.234	8.49	8.179	3.73	2.290
CD13	7.96	8.649	6.09	7.420	19.42	8.313
CD14	1.68	1.755	0.26	0.317	5.24	1.362
CD19	1.53	1.697	3.17	3.328	2.68	1.751
CD33	7.55	7.510	6.52	4.710	15.51	9.257
ALDH	68.96	14.630	47.19	11.882	1.40	0.500
CD35	1.57	1.239	0.48	0.424	9.58	8.285
CD38	68.75	16.872	69.92	20.245	17.66	3.871
CD41-C	27.22	18.522	25.35	19.138	26.32	14.980
CD45-M	54.62	18.533	51.03	22.604	70.51	13.339
CD56-BD	2.16	1.398	0.79	0.658	1.07	0.854
CD90	3.22	3.532	1.75	3.218	1.88	3.681
CD105	10.28	11.173	1.34	1.256	0.53	0.367
CD117	43.78	23.037	32.91	14.833	1.52	0.582
CD127	0.74	0.688	0.27	0.181	0.07	0.053
CD133	4.06	4.626	2.78	4.016	0.12	0.113
CD135	0.84	0.704	0.16	0.094	0.15	0.119
CD138	0.71	0.740	0.13	0.151	0.07	0.060
HLA-DR	38.43	16.437	49.76	18.578	14.92	2.690
GLY-A	5.67	4.942	0.38	0.228	9.16	5.722

Table 4A: Summary of Bone Marrow (All Gates)

Cord Blood								
CD Marker	ALDH CD+			CD34CD+			CD+	
	Mean	STDEV		Mean	STDEV		Mean	STDEV
CD3	1.95	1.249		3.15	1.863		43.69	14.643
CD7	7.36	14.051		6.30	6.090		43.76	17.460
CD10	0.46	0.410		1.07	1.334		0.79	0.847
CD13	16.33	21.495		14.60	21.476		22.12	10.275
CD14	1.13	1.359		0.25	0.188		22.62	3.440
CD19	0.55	0.660		1.53	0.872		6.45	3.990
CD33	12.33	13.595		9.70	12.607		23.73	3.985
ALDH	81.41	5.252		74.40	11.454		0.90	0.517
CD35	1.01	0.974		1.22	1.078		20.08	8.910
CD38	49.20	21.900		47.80	22.845		71.88	18.735
CD41-C	16.89	19.567		16.51	21.023		22.33	5.725
CD45-M	78.28	20.454		75.61	25.070		88.84	12.651
CD56-BD	4.64	9.899		3.17	7.043		5.90	7.961
CD90	8.00	14.188		6.14	11.781		6.24	10.195
CD105	0.82	0.686		0.23	0.187		1.44	2.131
CD117	45.96	26.423		45.36	25.923		0.67	0.500
CD127	0.39	0.255		0.14	0.159		0.59	1.012
CD133	29.61	21.858		30.74	24.161		0.32	0.255
CD135	1.09	1.428		0.34	0.401		0.34	0.463
CD138	0.11			0.09			0.37	
HLA-DR	65.79	20.136		64.26	22.024		31.27	6.338
GLY-A	1.46	0.479		0.85	0.556		3.07	2.110

Table 4B: Summary of Cord Blood (All Gates)

Peripheral Blood						
CD Marker	ALDH CD+		CD34CD+		CD+	
	Mean	STDEV	Mean	STDEV	Mean	STDEV
CD3	1.47	1.426	1.86	2.371	21.81	18.080
CD7	1.02	0.845	1.29	0.976	14.85	15.860
CD10	0.47	0.721	0.61	0.737	0.72	0.933
CD13	24.84	26.139	25.16	25.962	45.08	27.214
CD14	0.66	0.798	0.57	0.556	25.43	11.399
CD19	0.60	0.762	0.67	0.837	4.69	8.967
CD33	18.71	22.23	18.25	22.459	25.98	17.523
ALDH	94.85	4.405	83.41	11.946	0.98	1.082
CD35	2.37	2.65	2.06	2.040	41.67	21.568
CD38	69.17	26.779	67.65	27.161	36.31	18.805
CD41-C	27.26	23.166	26.94	23.251	41.81	19.781
CD45-M	86.08	21.323	84.27	23.342	96.93	4.692
CD56-BD	1.62	2.269	1.51	2.186	3.98	4.527
CD90	2.52	2.356	1.75	1.868	0.72	0.347
CD105	1.03	1.188	0.78	1.128	2.36	4.736
CD117	18.99	18.045	17.93	18.403	0.24	0.308
CD127	0.54	0.657	0.40	0.489	1.03	2.686
CD133	36.80	26.149	35.63	26.734	0.27	0.151
CD135	0.58	0.717	0.43	0.632	0.10	0.080
CD138	0.57	0.582	0.43	0.584	0.04	0.012
HLA-DR	64.89	26.688	62.44	26.427	21.49	10.799
GLY-A	0.69	0.694	0.51	0.611	0.47	0.379
ALDH	1.11	1.084	1.11	1.085	1.11	1.085

Table 4C: Summary of Peripheral Blood (All Gates)

Example 2: Visualization and Characterization of the ALDH^{br}SSC^{lo} Cells Having the Phenotype of Interest

For intracellular staining, FACS™ isolated ALDH^{br}SSC^{lo} cells having the phenotype of interest, as identified in Example 1, are stained using the Fix and Perm cell permeabilization kit according to the manufacturer's instructions (Caltag Laboratories). In preparation for differential Wright Giemsa staining, cells in PBS are pelleted for 3 minutes at 1000 rpm directly onto coated slides using a Cytospin3 centrifuge. The cells are stained with Wright Giemsa stain in an automated cell stainer.

Example 3: Functional Characterization of ALDH^{br}SSC^{lo} Cells Having the Phenotype of Interest Using *In Vitro* Culture

Hematopoietic progenitor colony HPC assays are performed by plating 100-200 ALDH^{br}SSC^{lo} cells having the phenotype of interest, as identified in Example 1, in MethoCult H4431 (StemCell Technologies, Inc.). The cells are incubated in a humidified chamber at 37°C with 5% CO₂. Hematopoietic colonies (>100 cells) are then scored at 14-18 days after initiating the cultures. See, for example, Fallon *et al.* (2003) *Br. J. Haematol.* 121:1, herein incorporated by reference in its entirety.

Long-term culture (LTC) assays are performed by maintaining ALDH^{br}SSC^{lo} cells having the phenotype of interest, as identified in Example 1, on either irradiated allogeneic bone marrow stroma or stromal layers of murine MS-5 cells (Issaad *et al.* (1993) *Blood* 81:2916) or in some instances S17 cells (with medium containing Flt3 ligand, IL-3, erythropoietin, G-CSF, and IL-15) according to the method of Fallon *et al.* (2003) *Br. J. Haematol.* 121:1. MS-5 cells support the growth of human multipotent cells and BFU-E for extended periods of time and may support early stem cells as well as, or better than, standard allogeneic stroma. The stromal layers are established by seeding the center wells of 24-well plates (Corning Costar Corp.) with 6-7x10⁴ MS-5 cells/well in 0.5 ml DMEM supplemented with 10% FCS. These cells are cultured at 37°C in a humidified incubator until the cultures approach approximately 80% confluence. The monolayers are then irradiated with 30 Gy γ -irradiation from a cesium source. After irradiation, the culture media from the monolayers is replaced entirely with Myelocult

H5100 (StemCell Technologies, Inc.) and the cells are maintained at 33°C in a humidified chamber with 5% CO₂. Long-term cultures are typically initiated with 500-20,000 sorted hematopoietic progenitor cells/well on the irradiated MS-5 cells. At 7-10 day intervals, half the media from each well is removed so that the media can be replenished. Adherent and non-adherent cells are harvested after 2 weeks and plated into HPC assays as described above. The HPC assay generally identifies relatively mature progenitor cells with limited lineage and self-renewal potential, while the LTC assay generally quantifies more primitive cells with a higher self-renewal potential.

Example 4: Large Segmental Canine Femoral Defects Are Healed with Autologous Stem Cell Therapy

The stem cells of the invention are used in a canine model of bone engraftment. See, for example, U.S. Patent No. 6,541,024 herein incorporated by reference in its entirety. The following protocols are used for culture-expanding autologous mesenchymal stem cells of the invention that can regenerate clinically significant bone defects in a large animal model.

MSC Cultivation and Manipulation

A 15 cc bone marrow aspirate is obtained from the iliac crest of each animal, according to an IACUC-approved protocol, and shipped on ice by overnight courier to the cell culture facilities. Isolation of canine ALDH^{br} MSCs is achieved using procedures described in Example 1. Tissue culture flasks (185 cm²) are seeded with 10⁷ nucleated cells isolated from the cushion, and cultured with DMEM containing 10% fetal calf serum from a selected lot. Cells are passaged at 8x10³ cells/cm², and maintained until the time of implantation. Cell-loaded implants are prepared by incubating fibronectin-coated porous hydroxyapatite-tricalcium phosphate (HA/TCP) cylinders (Zimmer, Inc.) in a 7.5x10⁶ cells/ml suspension of MSCs for 3 hr at 37° C. The interval between marrow harvest and implantation is 16 days. An aliquot of cells from each preparation is also cultured under osteoinductive conditions to quantify aspects of osteoblastic differentiation.

Canine Femoral Gap Model

A unilateral segmental femoral defect model is developed for this study following IACUC approval. Under general anesthesia, thirty-six skeletally mature female purpose-bred hounds (20 kg) undergo resection of a 21 mm long osteoperiosteal segment from their mid-diaphysis. A 4.5 mm Synthes™ 8-hole lengthening plate is contoured to the lateral aspect of the bone, and secured with bicortical screws. The defect is filled with one of three materials; 1) a cell-free HA/TCP cylinder, 2) an MSC-loaded HA/TCP cylinder, or 3) cancellous bone harvested from the iliac crest. HA/TCP implants are secured by placing two sutures around the implant and the plate. Animals receive peri-operative antibiotics, and analgesics are administered for three days post-operatively.

Radiographic and Histologic Analyses

Standard radiographic images are obtained at pre-op, immediately post-op, and at 4 week intervals until termination of the study. All samples contain a radiodensity step wedge to provide a basis for comparing changes over time, and between dogs. Upon sacrifice, specimens are subjected to high resolution Faxitron radiography, and subsequently processed for biomechanical evaluation. Following torsion testing, undecalcified longitudinal sections will be processed for quantitative histomorphometry.

Biomechanical Testing

Sixteen weeks after implantation, animals are sacrificed for torsion testing of femurs. The fixation plate, screws, and adherent soft tissue are removed, and the metaphyses of the bones are embedded. The specimens are externally rotated in a custom torsion test apparatus, failure load and stiffness recorded, and the data analyzed by one way ANOVA according to post hoc Student-Newman-Keuls tests.

Results

Generally, animals tolerate the surgical procedure well, with no incidence of infection, implant rejection, or failure of fixation. Two modes of repair are generally observable in the MSC-loaded samples; first, considerable callus formation generally occurs at both host-implant interfaces; and second, a substantial collar of bone

surrounding the implant itself develops. Cell-free implants generally do not possess either of these features. Autograft samples generally undergo a traditional consolidation sequence, with the majority of bone being laid down in the medial aspect of the gap defect. Generally, MSC-loaded samples not only became fully integrated at the host implant interface, but the periosteal collar extended proximally and distally beyond the cut edges of the gap. Furthermore, the diameter of new bone at the mid-diaphysis is generally greater in MSC-loaded implants than either autograft samples or intact limbs. Analyses of the osteogenic potential MSCs from each animal generally demonstrate the development of alkaline phosphatase-positive cells which deposit significant mineralized extracellular matrix.

Many modifications and other embodiments of the inventions set forth herein will come to mind to one skilled in the art to which these inventions pertain having the benefit of the teachings presented in the foregoing descriptions and the associated drawings. Therefore, it is to be understood that the inventions are not to be limited to the specific embodiments disclosed and that modifications and other embodiments are intended to be included within the scope of the appended claims. Although specific terms are employed herein, they are used in a generic and descriptive sense only and not for purposes of limitation.

All publications and patent applications mentioned in the specification are indicative of the level of those skilled in the art to which this invention pertains. All publications and patent applications are herein incorporated by reference to the same extent as if each individual publication or patent application was specifically and individually indicated to be incorporated by reference.

THAT WHICH IS CLAIMED:

1. A cell population of ALDH^{br}SSC^{lo} stem cells, wherein greater than about 60% of the cells within the population express the cell surface marker CD105 wherein said cells are capable of multilineage development.

STEM CELL POPULATIONS AND METHODS OF USE

ABSTRACT OF THE DISCLOSURE

The present invention relates to populations of stem cells, methods for isolating these stem cell populations, and methods repairing, regenerating , and reconstituting tissues using the these stem cell populations. The invention additionally relates to methods of screening agents that promote growth, engraftment, or differentiation of stem cells.

Compensation in Multiparameter Analysis of ALDH^{br}SSC^l Cells

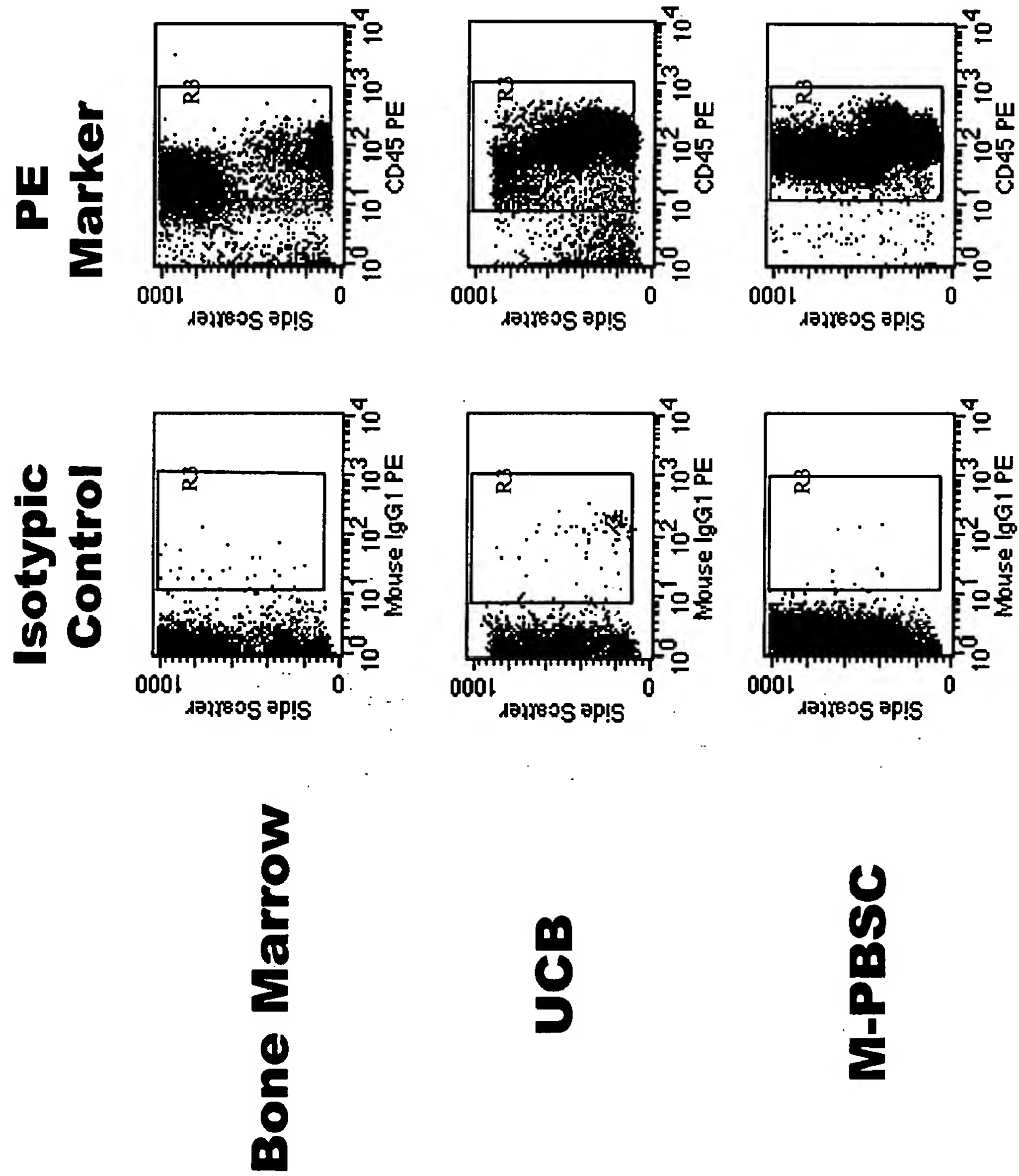


Fig. 1

Expression of Surface Markers in Different Graft : All Cells

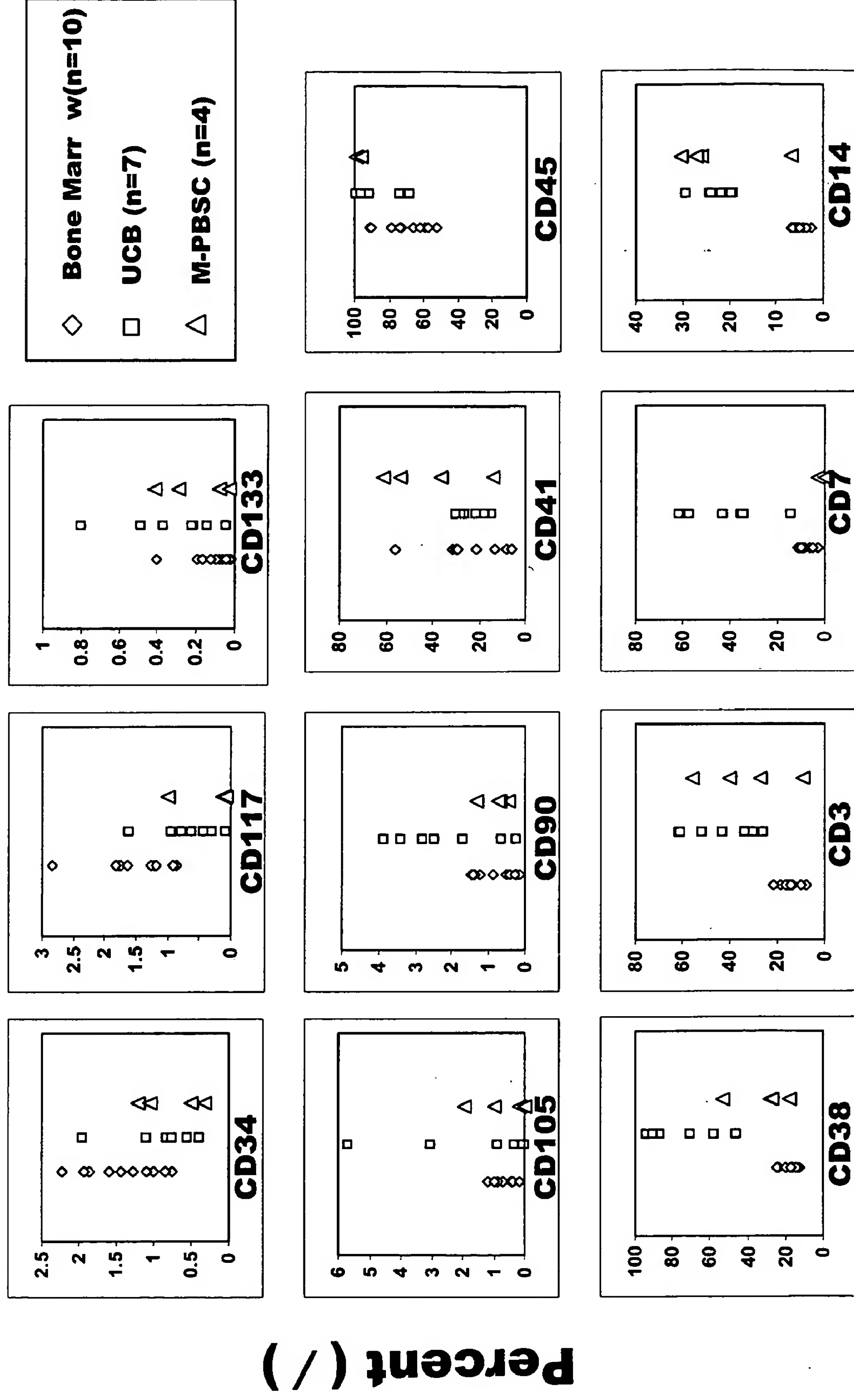


Fig. 2

Expression of Stem/Progenitor Markers in Different Graft : **ALDH^{br}SSC^{lo} Population**

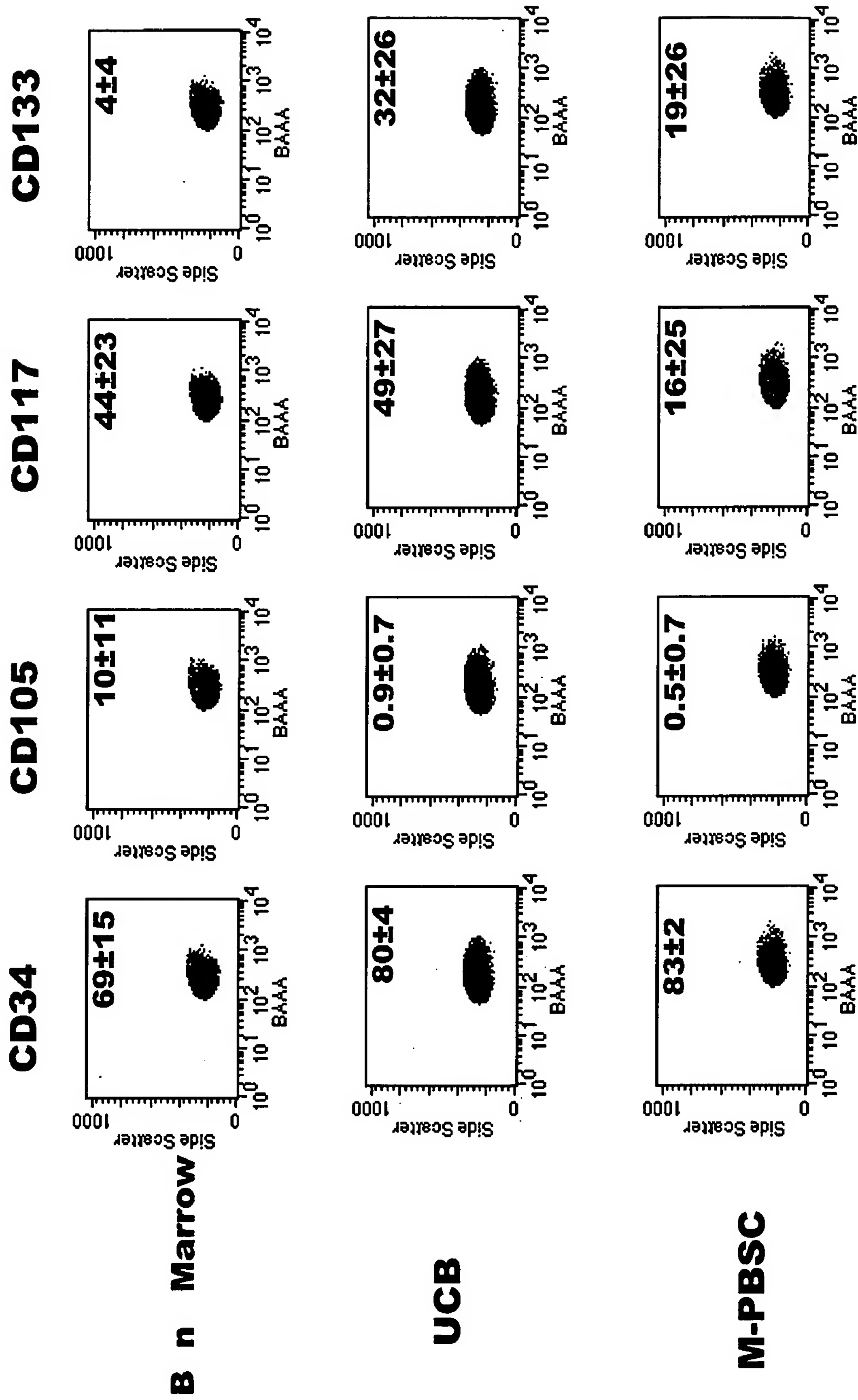


Fig. 3

Expression of Surface Markers in Different Grafts: ALDH^{br}SSC^{lo} Population

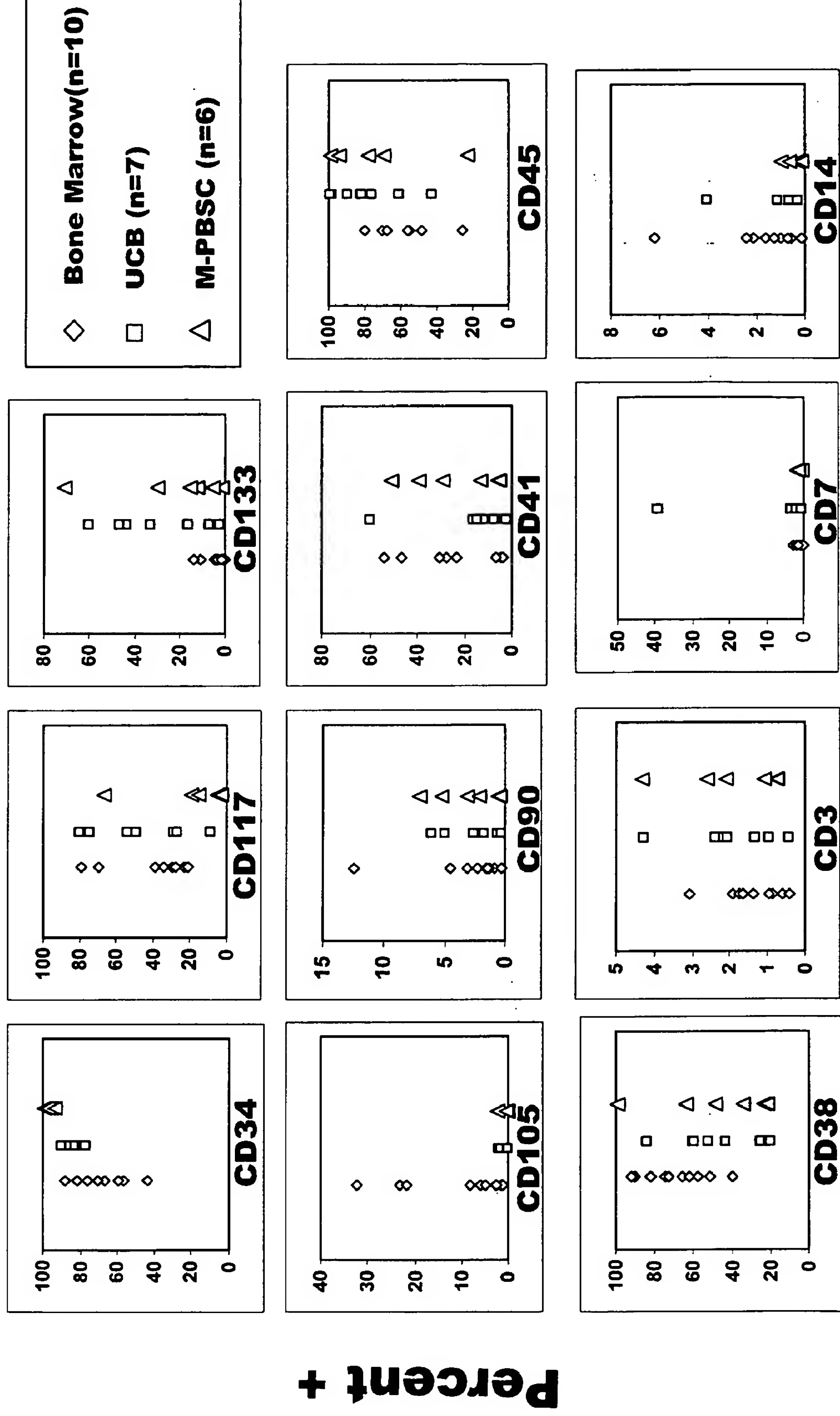


Fig. 4

ALDH^{br} and ALDH^{dim} Subpopulations **7AAD negative**

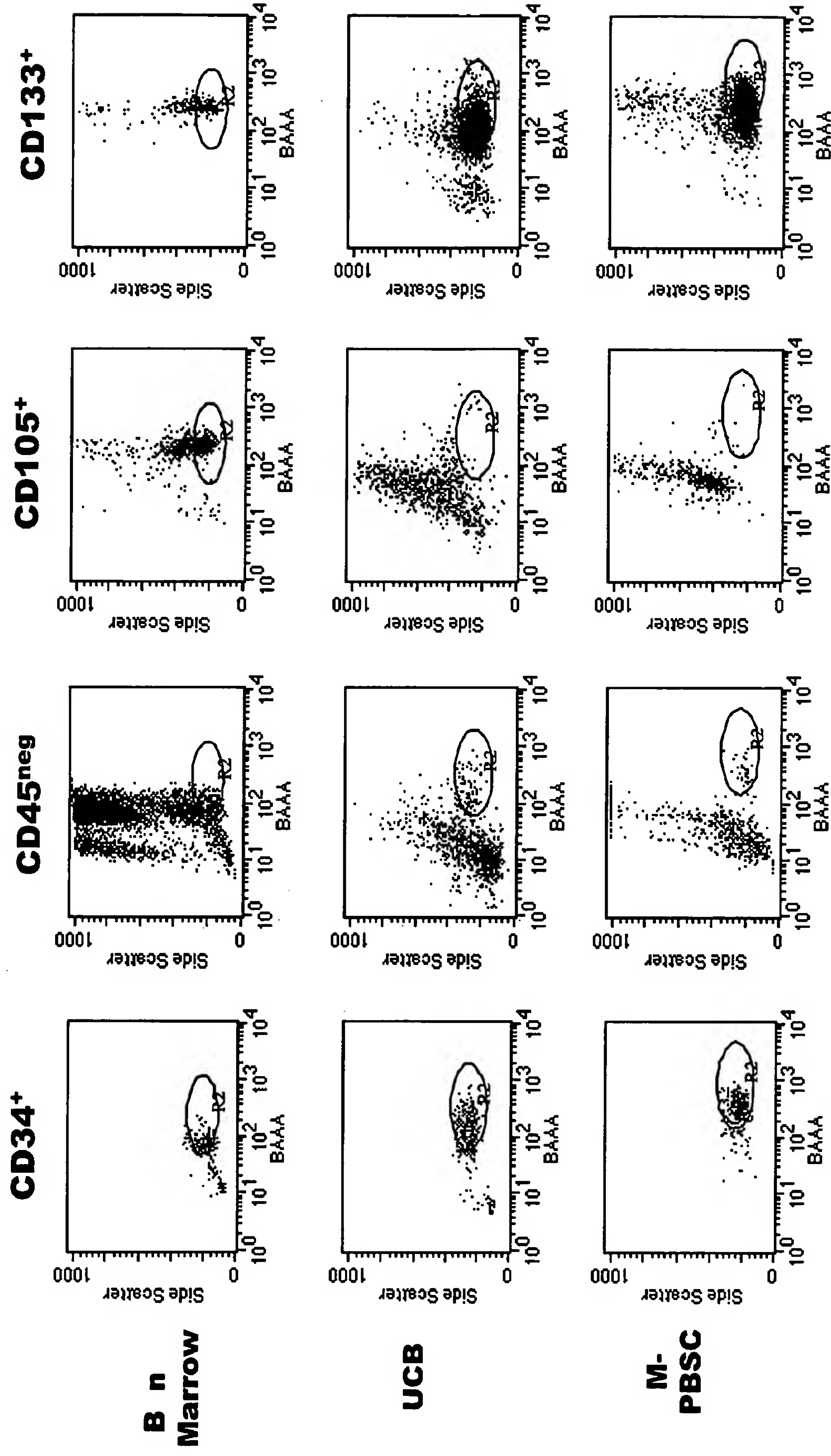


Fig. 5

Comparison of CD34 and ALDH^{br} Subpopulations

Bon Marrow

	ALDH ^{br} SSC ^{lo} Marker ⁺		CD34 ⁺ Marker ⁺	
Marker	Mean	SD	Mean	SD
CD38	68.75	16.87	69.92	20.25
CD45-M	54.62	18.53	51.03	22.60
CD105	10.28	11.17	1.34	1.26
CD133	4.06	4.63	2.78	4.02

UCB

	ALDH ^{br} SSC ^{lo} Marker ⁺		CD34 ⁺ Marker ⁺	
Marker	Mean	SD	Mean	SD
CD38	49.20	21.90	47.80	22.84
CD45-M	78.28	20.45	75.61	25.07
CD105	0.82	0.69	0.23	0.19
CD133	29.61	21.86	30.74	24.16

M-PBSC

	ALDH ^{br} SSC ^{lo} Marker ⁺		CD34 ⁺ Marker ⁺	
Marker	Mean	SD	Mean	SD
CD38	69.17	26.78	67.65	27.16
CD45-M	86.08	21.32	84.27	23.34
CD105	1.03	1.19	0.78	1.13
CD133	36.80	26.15	35.63	26.73

Fig. 6